

IN THE UNITED STATES PATENT AND TRADEMARK OFFICE

PATENT

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DIAGNOSIS OF LUNG CANCER



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Washington, DC 20231

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Respectfully submitted,

Seed Intellectual Property Law Group PLLC

Steve Plante/Jeanette West/Susan Johnson

JEP:sds

Enclosures:

- Postcard
- Form PTO/SB/05
- Specification, Claims, Abstract (187 pages)
- Sequence Listing (518 pages)
- Declaration for Sequence Listing
- Diskette for Sequence Listing

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COMPOSITIONS AND METHODS FOR THE THERAPY AND DIAGNOSIS OF LUNG CANCER

REFERENCE TO RELATED APPLICATIONS

This application is a continuation-in-part of U.S. Patent Application
5 No. 09/614,124, filed July 11, 2000, which is a continuation-in-part of U.S. Patent
Application No. 09/589,184, filed June 5, 2000, which is a continuation-in-part of U.S.
Patent Application No. 09/560,406, filed April 27, 2000, which is a continuation-in-part of
U.S. Patent Application No. 09/546,259, filed April 10, 2000, which is a continuation-in-
part of U.S. Patent Application No. 09/533,077, filed March 22, 2000, which is a
10 continuation-in-part of U.S. Patent Application No. 09/519,642 filed March 6, 2000, which
is a continuation-in-part of U.S. Patent Application No. 09/476,300, filed December 30,
1999, which is a continuation-in-part of U.S. Patent Application No. 09/466,867, filed
December 17, 1999, which is a continuation-in-part of U.S. Patent Application 09/419,356,
filed October 15, 1999, which is a continuation-in-part of U.S. Patent Application
15 No. 09/346,492, filed June 30, 1999, and is related to PCT/US00/18061, filed 6/30/00.

TECHNICAL FIELD OF THE INVENTION

The present invention relates generally to therapy and diagnosis of cancer,
such as lung cancer. The invention is more specifically related to polypeptides comprising
at least a portion of a lung tumor protein, and to polynucleotides encoding such
20 polypeptides. Such polypeptides and polynucleotides may be used in compositions for
prevention and treatment of lung cancer, and for the diagnosis and monitoring of such
cancers.

BACKGROUND OF THE INVENTION

Cancer is a significant health problem throughout the world. Although
25 advances have been made in detection and therapy of cancer, no vaccine or other
universally successful method for prevention or treatment is currently available. Current

therapies, which are generally based on a combination of chemotherapy or surgery and radiation, continue to prove inadequate in many patients.

Lung cancer is the primary cause of cancer death among both men and women in the U.S., with an estimated 172,000 new cases being reported in 1994. The five-
 5 year survival rate among all lung cancer patients, regardless of the stage of disease at diagnosis, is only 13%. This contrasts with a five-year survival rate of 46% among cases detected while the disease is still localized. However, only 16% of lung cancers are discovered before the disease has spread.

Early detection is difficult since clinical symptoms are often not seen until
 10 the disease has reached an advanced stage. Currently, diagnosis is aided by the use of chest x-rays, analysis of the type of cells contained in sputum and fiberoptic examination of the bronchial passages. Treatment regimens are determined by the type and stage of the cancer, and include surgery, radiation therapy and/or chemotherapy.

In spite of considerable research into therapies for this and other cancers,
 15 lung cancer remains difficult to diagnose and treat effectively. Accordingly, there is a need in the art for improved methods for detecting and treating such cancers. The present invention fulfills these needs and further provides other related advantages.

SUMMARY OF THE INVENTION

Briefly stated, the present invention provides compositions and methods for
 20 the diagnosis and therapy of cancer, such as lung cancer. In one aspect, the present invention provides polypeptides comprising at least a portion of a lung tumor protein, or a variant thereof. Certain portions and other variants are immunogenic, such that the ability of the variant to react with antigen-specific antisera is not substantially diminished. Within certain embodiments, the polypeptide comprises a sequence that is encoded by a
 25 polynucleotide sequence selected from the group consisting of: (a) sequences recited in SEQ ID NO: 1, 11-13, 15, 20, 23-27, 29, 30, 33, 34, 39, 41, 43-46, 51, 52, 57, 58, 60, 62, 65-67, 69-71, 74, 76, 79, 80, 84, 86, 89-92, 95, 97, 98, 101, 110, 111, 113-119, 121-128, 130-134, 136, 138, 139, 141, 143, 146-151, 153, 154, 157-160, 162-164, 167-178, 180,

181, 183, 186-190, 192, 193, 195-220, 224, 226-231, 234, 236, 237, 240, 241, 244-246, 248, 254, 255, 261, 262, 266, 270, 275, 280, 282, 283, 288, 289, 290, 292, 295, 301, 303, 304, 309, 311, 341-782, 784, 785, 790, 792, 794, 796, 800-804, 807, 808 and 810-826, 1240, 1243, 1247, 1269, 1272, 1280, 1283, 1285, 1286, 1289, 1300, 1309, 1318, 1319, 5 1327, 1335, 1339, 1346, 1359, 1369, 1370, 1371, 1393, 1398, 1405, 1408, 1413, 1414, 1417, 1422, 1429, 1432, 1435, 1436, 1438-1442, 1447, 1450, 1453, 1463, 1467, 1470, 1473, 1475, 1482, 1486, 1491-1494, 1501, 1505, 1506, 1514-1517, 1520, 1522, 1524, 1535, 1538, 1542, 1543, 1547, 1554, 1557, 1559, 1561, 1563, and 1669; (b) variants of a sequence recited in SEQ ID NO: 1, 11-13, 15, 20, 23-27, 29, 30, 33, 34, 39, 41, 43-46, 51, 10 52, 57, 58, 60, 62, 65-67, 69-71, 74, 76, 79, 80, 84, 86, 89-92, 95, 97, 98, 101, 110, 111, 113-119, 121-128, 130-134, 136, 138, 139, 141, 143, 146-151, 153, 154, 157-160, 162-164, 167-178, 180, 181, 183, 186-190, 192, 193, 195-220, 224, 226-231, 234, 236, 237, 240, 241, 244-246, 248, 254, 255, 261, 262, 266, 270, 275, 280, 282, 283, 288, 289, 290, 292, 295, 301, 303, 304, 309, 311, 341-782, 784, 785, 790, 792, 794, 796, 800-804, 807, 15 808, 810-826, 1240, 1243, 1247, 1269, 1272, 1280, 1283, 1285, 1286, 1289, 1300, 1309, 1318, 1319, 1327, 1335, 1339, 1346, 1359, 1369, 1370, 1371, 1393, 1398, 1405, 1408, 1413, 1414, 1417, 1422, 1429, 1432, 1435, 1436, 1438-1442, 1447, 1450, 1453, 1463, 1467, 1470, 1473, 1475, 1482, 1486, 1491-1494, 1501, 1505, 1506, 1514-1517, 1520, 1522, 1524, 1535, 1538, 1542, 1543, 1547, 1554, 1557, 1559, 1561, 1563, and 1669; and 20 (c) complements of a sequence of (a) or (b). In specific embodiments, the polypeptides of the present invention comprise at least a portion of a tumor protein that includes an amino acid sequence selected from the group consisting of sequences recited in SEQ ID NO: 786, 787, 791, 793, 795, 797-799, 806, 809, 827, 1670-1675 and 1677-1678 and variants thereof.

25 The present invention further provides polynucleotides that encode a polypeptide as described above, or a portion thereof (such as a portion encoding at least 15 amino acid residues of a lung tumor protein), expression vectors comprising such polynucleotides and host cells transformed or transfected with such expression vectors.

Within other aspects, the present invention provides pharmaceutical compositions comprising a polypeptide or polynucleotide as described above and a physiologically acceptable carrier.

5 Within a related aspect of the present invention, vaccines, or immunogenic compositions, for prophylactic or therapeutic use are provided. Such vaccines comprise a polypeptide or polynucleotide as described above and an immunostimulant.

The present invention further provides pharmaceutical compositions that comprise: (a) an antibody or antigen-binding fragment thereof that specifically binds to a lung tumor protein; and (b) a physiologically acceptable carrier.

10 Within further aspects, the present invention provides pharmaceutical compositions comprising: (a) an antigen presenting cell that expresses a polypeptide as described above and (b) a pharmaceutically acceptable carrier or excipient. Antigen presenting cells include dendritic cells, macrophages, monocytes, fibroblasts and B cells.

15 Within related aspects, vaccines, or immunogenic compositions, are provided that comprise: (a) an antigen presenting cell that expresses a polypeptide as described above and (b) an immunostimulant.

The present invention further provides, in other aspects, fusion proteins that comprise at least one polypeptide as described above, as well as polynucleotides encoding such fusion proteins.

20 Within related aspects, pharmaceutical compositions comprising a fusion protein, or a polynucleotide encoding a fusion protein, in combination with a physiologically acceptable carrier are provided.

25 Vaccines, or immunogenic compositions, are further provided, within other aspects, that comprise a fusion protein, or a polynucleotide encoding a fusion protein, in combination with an immunostimulant.

Within further aspects, the present invention provides methods for inhibiting the development of a cancer in a patient, comprising administering to a patient a pharmaceutical composition or immunogenic composition as recited above. The patient

may be afflicted with lung cancer, in which case the methods provide treatment for the disease, or patient considered at risk for such a disease may be treated prophylactically.

The present invention further provides, within other aspects, methods for removing tumor cells from a biological sample, comprising contacting a biological sample
5 with T cells that specifically react with a lung tumor protein, wherein the step of contacting is performed under conditions and for a time sufficient to permit the removal of cells expressing the protein from the sample.

Within related aspects, methods are provided for inhibiting the development of a cancer in a patient, comprising administering to a patient a biological sample treated as
10 described above.

Methods are further provided, within other aspects, for stimulating and/or expanding T cells specific for a lung tumor protein, comprising contacting T cells with one or more of: (i) a polypeptide as described above; (ii) a polynucleotide encoding such a polypeptide; and/or (iii) an antigen presenting cell that expresses such a polypeptide; under
15 conditions and for a time sufficient to permit the stimulation and/or expansion of T cells. Isolated T cell populations comprising T cells prepared as described above are also provided.

Within further aspects, the present invention provides methods for inhibiting the development of a cancer in a patient, comprising administering to a patient an effective
20 amount of a T cell population as described above.

The present invention further provides methods for inhibiting the development of a cancer in a patient, comprising the steps of: (a) incubating CD4⁺ and/or CD8⁺ T cells isolated from a patient with one or more of: (i) a polypeptide comprising at least an immunogenic portion of a lung tumor protein; (ii) a polynucleotide encoding such
25 a polypeptide; and (iii) an antigen-presenting cell that expressed such a polypeptide; and (b) administering to the patient an effective amount of the proliferated T cells, and thereby inhibiting the development of a cancer in the patient. Proliferated cells may, but need not, be cloned prior to administration to the patient.

Within further aspects, the present invention provides methods for determining the presence or absence of a cancer in a patient, comprising: (a) contacting a biological sample obtained from a patient with a binding agent that binds to a polypeptide as recited above; (b) detecting in the sample an amount of polypeptide that binds to the binding agent; and (c) comparing the amount of polypeptide with a predetermined cut-off value, and therefrom determining the presence or absence of a cancer in the patient. Within preferred embodiments, the binding agent is an antibody, more preferably a monoclonal antibody. The cancer may be lung cancer.

The present invention also provides, within other aspects, methods for monitoring the progression of a cancer in a patient. Such methods comprise the steps of: (a) contacting a biological sample obtained from a patient at a first point in time with a binding agent that binds to a polypeptide as recited above; (b) detecting in the sample an amount of polypeptide that binds to the binding agent; (c) repeating steps (a) and (b) using a biological sample obtained from the patient at a subsequent point in time; and (d) comparing the amount of polypeptide detected in step (c) with the amount detected in step (b) and therefrom monitoring the progression of the cancer in the patient.

The present invention further provides, within other aspects, methods for determining the presence or absence of a cancer in a patient, comprising the steps of: (a) contacting a biological sample obtained from a patient with an oligonucleotide that hybridizes to a polynucleotide that encodes a lung tumor protein; (b) detecting in the sample a level of a polynucleotide, preferably mRNA, that hybridizes to the oligonucleotide; and (c) comparing the level of polynucleotide that hybridizes to the oligonucleotide with a predetermined cut-off value, and therefrom determining the presence or absence of a cancer in the patient. Within certain embodiments, the amount of mRNA is detected via polymerase chain reaction using, for example, at least one oligonucleotide primer that hybridizes to a polynucleotide encoding a polypeptide as recited above, or a complement of such a polynucleotide. Within other embodiments, the amount of mRNA is detected using a hybridization technique, employing an

oligonucleotide probe that hybridizes to a polynucleotide that encodes a polypeptide as recited above, or a complement of such a polynucleotide.

In related aspects, methods are provided for monitoring the progression of a cancer in a patient, comprising the steps of: (a) contacting a biological sample obtained from a patient with an oligonucleotide that hybridizes to a polynucleotide that encodes a lung tumor protein; (b) detecting in the sample an amount of a polynucleotide that hybridizes to the oligonucleotide; (c) repeating steps (a) and (b) using a biological sample obtained from the patient at a subsequent point in time; and (d) comparing the amount of polynucleotide detected in step (c) with the amount detected in step (b) and therefrom monitoring the progression of the cancer in the patient.

Within further aspects, the present invention provides antibodies, such as monoclonal antibodies, that bind to a polypeptide as described above, as well as diagnostic kits comprising such antibodies. Diagnostic kits comprising one or more oligonucleotide probes or primers as described above are also provided.

These and other aspects of the present invention will become apparent upon reference to the following detailed description. All references disclosed herein are hereby incorporated by reference in their entirety as if each was incorporated individually.

SEQUENCE IDENTIFIERS

SEQ ID NO: 1 is the determined cDNA sequence for clone #19038, also referred to as L845P.

SEQ ID NO: 2 is the determined cDNA sequence for clone #19036.

SEQ ID NO: 3 is the determined cDNA sequence for clone #19034.

SEQ ID NO: 4 is the determined cDNA sequence for clone #19033.

SEQ ID NO: 5 is the determined cDNA sequence for clone #19032.

SEQ ID NO: 6 is the determined cDNA sequence for clone #19030, also referred to as L559S.

SEQ ID NO: 7 is the determined cDNA sequence for clone #19029.

SEQ ID NO: 8 is the determined cDNA sequence for clone #19025.

SEQ ID NO: 9 is the determined cDNA sequence for clone #19023.

SEQ ID NO: 10 is the determined cDNA sequence for clone #18929.

SEQ ID NO: 11 is the determined cDNA sequence for clone #19010.

SEQ ID NO: 12 is the determined cDNA sequence for clone #19009.

5 SEQ ID NO: 13 is the determined cDNA sequence for clones #19005, 19007, 19016 and 19017.

SEQ ID NO: 14 is the determined cDNA sequence for clone #19004.

SEQ ID NO: 15 is the determined cDNA sequence for clones #19002 and 18965.

10 SEQ ID NO: 16 is the determined cDNA sequence for clone #18998.

SEQ ID NO: 17 is the determined cDNA sequence for clone #18997.

SEQ ID NO: 18 is the determined cDNA sequence for clone #18996.

SEQ ID NO: 19 is the determined cDNA sequence for clone #18995.

15 SEQ ID NO: 20 is the determined cDNA sequence for clone #18994, also known as L846P.

SEQ ID NO: 21 is the determined cDNA sequence for clone #18992.

SEQ ID NO: 22 is the determined cDNA sequence for clone #18991.

SEQ ID NO: 23 is the determined cDNA sequence for clone #18990, also referred to as clone #20111.

20 SEQ ID NO: 24 is the determined cDNA sequence for clone #18987.

SEQ ID NO: 25 is the determined cDNA sequence for clone #18985, also referred as L839P.

SEQ ID NO: 26 is the determined cDNA sequence for clone #18984, also referred to as L847P.

25 SEQ ID NO: 27 is the determined cDNA sequence for clone #18983.

SEQ ID NO: 28 is the determined cDNA sequence for clones #18976 and 18980.

SEQ ID NO: 29 is the determined cDNA sequence for clone #18975.

SEQ ID NO: 30 is the determined cDNA sequence for clone #18974.

SEQ ID NO: 31 is the determined cDNA sequence for clone #18973.

SEQ ID NO: 32 is the determined cDNA sequence for clone #18972.

SEQ ID NO: 33 is the determined cDNA sequence for clone #18971, also referred to as L801P.

5 SEQ ID NO: 34 is the determined cDNA sequence for clone #18970.

SEQ ID NO: 35 is the determined cDNA sequence for clone #18966.

SEQ ID NO: 36 is the determined cDNA sequence for clones #18964, 18968 and 19039.

SEQ ID NO: 37 is the determined cDNA sequence for clone #18960.

10 SEQ ID NO: 38 is the determined cDNA sequence for clone #18959.

SEQ ID NO: 39 is the determined cDNA sequence for clones #18958 and 18982.

SEQ ID NO: 40 is the determined cDNA sequence for clones #18956 and 19015.

15 SEQ ID NO: 41 is the determined cDNA sequence for clone #18954, also referred to L848P.

SEQ ID NO: 42 is the determined cDNA sequence for clone #18951.

SEQ ID NO: 43 is the determined cDNA sequence for clone #18950.

20 SEQ ID NO: 44 is the determined cDNA sequence for clones #18949 and 19024, also referred to as L844P.

SEQ ID NO: 45 is the determined cDNA sequence for clone #18948.

SEQ ID NO: 46 is the determined cDNA sequence for clone #18947, also referred to as L840P.

25 SEQ ID NO: 47 is the determined cDNA sequence for clones #18946, 18953, 18969 and 19027.

SEQ ID NO: 48 is the determined cDNA sequence for clone #18942.

SEQ ID NO: 49 is the determined cDNA sequence for clone #18940, 18962, 18963, 19006, 19008, 19000, and 19031.

SEQ ID NO: 50 is the determined cDNA sequence for clone #18939.

SEQ ID NO: 51 is the determined cDNA sequence for clones #18938 and 18952.

SEQ ID NO: 52 is the determined cDNA sequence for clone #18938.

SEQ ID NO: 53 is the determined cDNA sequence for clone #18937.

5 SEQ ID NO: 54 is the determined cDNA sequence for clones #18934, 18935, 18993 and 19022, also referred to as L548S.

SEQ ID NO: 55 is the determined cDNA sequence for clone #18932.

SEQ ID NO: 56 is the determined cDNA sequence for clones #18931 and 18936.

10 SEQ ID NO: 57 is the determined cDNA sequence for clone #18930.

SEQ ID NO: 58 is the determined cDNA sequence for clone #19014, also referred to as L773P.

SEQ ID NO: 59 is the determined cDNA sequence for clone #19127.

15 SEQ ID NO: 60 is the determined cDNA sequence for clones #19057 and 19064.

SEQ ID NO: 61 is the determined cDNA sequence for clone #19122.

SEQ ID NO: 62 is the determined cDNA sequence for clones #19120 and 18121.

SEQ ID NO: 63 is the determined cDNA sequence for clone #19118.

20 SEQ ID NO: 64 is the determined cDNA sequence for clone #19117.

SEQ ID NO: 65 is the determined cDNA sequence for clone #19116.

SEQ ID NO: 66 is the determined cDNA sequence for clone #19114.

SEQ ID NO: 67 is the determined cDNA sequence for clone #19112, also known as L561S.

25 SEQ ID NO: 68 is the determined cDNA sequence for clone #19110.

SEQ ID NO: 69 is the determined cDNA sequence for clone #19107, also referred to as L552S.

SEQ ID NO: 70 is the determined cDNA sequence for clone #19106, also referred to as L547S.

SEQ ID NO: 71 is the determined cDNA sequence for clones #19105 and 19111.

SEQ ID NO: 72 is the determined cDNA sequence for clone #19099.

SEQ ID NO: 73 is the determined cDNA sequence for clones #19095,
5 19104 and 19125, also referred to as L549S.

SEQ ID NO: 74 is the determined cDNA sequence for clone #19094.

SEQ ID NO: 75 is the determined cDNA sequence for clones #19089 and 19101.

SEQ ID NO: 76 is the determined cDNA sequence for clone #19088.

10 SEQ ID NO: 77 is the determined cDNA sequence for clones #19087, 19092, 19096, 19100 and 19119.

SEQ ID NO: 78 is the determined cDNA sequence for clone #19086.

SEQ ID NO: 79 is the determined cDNA sequence for clone #19085, also referred to as L550S.

15 SEQ ID NO: 80 is the determined cDNA sequence for clone #19084, also referred to as clone #19079.

SEQ ID NO: 81 is the determined cDNA sequence for clone #19082.

SEQ ID NO: 82 is the determined cDNA sequence for clone #19080.

SEQ ID NO: 83 is the determined cDNA sequence for clone #19077.

20 SEQ ID NO: 84 is the determined cDNA sequence for clone #19076, also referred to as L551S.

SEQ ID NO: 85 is the determined cDNA sequence for clone #19074, also referred to as clone #20102.

25 SEQ ID NO: 86 is the determined cDNA sequence for clone #19073, also referred to as L560S.

SEQ ID NO: 87 is the determined cDNA sequence for clones #19072 and 19115.

SEQ ID NO: 88 is the determined cDNA sequence for clone #19071.

SEQ ID NO: 89 is the determined cDNA sequence for clone #19070.

SEQ ID NO: 90 is the determined cDNA sequence for clone #19069.

SEQ ID NO: 91 is the determined cDNA sequence for clone #19068, also referred to L563S.

SEQ ID NO: 92 is the determined cDNA sequence for clone #19066.

5 SEQ ID NO: 93 is the determined cDNA sequence for clone #19065.

SEQ ID NO: 94 is the determined cDNA sequence for clone #19063.

SEQ ID NO: 95 is the determined cDNA sequence for clones #19061, 19081, 19108 and 19109.

10 SEQ ID NO: 96 is the determined cDNA sequence for clones #19060, 19067 and 19083, also referred to as L548S.

SEQ ID NO: 97 is the determined cDNA sequence for clones #19059 and 19062.

SEQ ID NO: 98 is the determined cDNA sequence for clone #19058.

SEQ ID NO: 99 is the determined cDNA sequence for clone #19124.

15 SEQ ID NO: 100 is the determined cDNA sequence for clone #18929.

SEQ ID NO: 101 is the determined cDNA sequence for clone #18422.

SEQ ID NO: 102 is the determined cDNA sequence for clone #18425.

SEQ ID NO: 103 is the determined cDNA sequence for clone #18431.

SEQ ID NO: 104 is the determined cDNA sequence for clone #18433.

20 SEQ ID NO: 105 is the determined cDNA sequence for clone #18444.

SEQ ID NO: 106 is the determined cDNA sequence for clone #18449.

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SEQ ID NO: 109 is the determined cDNA sequence for clone #18455.

25 SEQ ID NO: 110 is the determined cDNA sequence for clone #18457.

SEQ ID NO: 111 is the determined cDNA sequence for clone #18466.

SEQ ID NO: 112 is the determined cDNA sequence for clone #18468.

SEQ ID NO: 113 is the determined cDNA sequence for clone #18471.

SEQ ID NO: 114 is the determined cDNA sequence for clone #18475.

SEQ ID NO: 115 is the determined cDNA sequence for clone #18476.
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 10 SEQ ID NO: 124 is the determined cDNA sequence for clone #20653.
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 25 SEQ ID NO: 165 is the determined cDNA sequence for clone #19146.
 SEQ ID NO: 166 is the determined cDNA sequence for clone #19149.1.
 SEQ ID NO: 167 is the determined cDNA sequence for clone #19152.
 SEQ ID NO: 168 is a first determined cDNA sequence for clone #19153.1.

SEQ ID NO: 169 is a second determined cDNA sequence for clone #19153.2.

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SEQ ID NO: 171 is the determined cDNA sequence for clone #19157.

5 SEQ ID NO: 172 is the determined cDNA sequence for clone #19159.

SEQ ID NO: 173 is the determined cDNA sequence for clone #19160.

SEQ ID NO: 174 is a first determined cDNA sequence for clone #19161.1.

SEQ ID NO: 175 is a second determined cDNA sequence for clone #19161.2.

10 SEQ ID NO: 176 is the determined cDNA sequence for clone #19162.1.

SEQ ID NO: 177 is the determined cDNA sequence for clone #19166.

SEQ ID NO: 178 is the determined cDNA sequence for clone #19169.

SEQ ID NO: 179 is the determined cDNA sequence for clone #19171.

SEQ ID NO: 180 is a first determined cDNA sequence for clone #19173.1.

15 SEQ ID NO: 181 is a second determined cDNA sequence for clone #19173.2.

SEQ ID NO: 182 is the determined cDNA sequence for clone #19174.1.

SEQ ID NO: 183 is the determined cDNA sequence for clone #19175.

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20 SEQ ID NO: 185 is the determined cDNA sequence for clone #19178.

SEQ ID NO: 186 is the determined cDNA sequence for clone #19179.1.

SEQ ID NO: 187 is the determined cDNA sequence for clone #19179.2.

SEQ ID NO: 188 is the determined cDNA sequence for clone #19180.

SEQ ID NO: 189 is a first determined cDNA sequence for clone #19182.1.

25 SEQ ID NO: 190 is a second determined cDNA sequence for clone #19182.2.

SEQ ID NO: 191 is the determined cDNA sequence for clone #19183.1.

SEQ ID NO: 192 is the determined cDNA sequence for clone #19185.1.

SEQ ID NO: 193 is the determined cDNA sequence for clone #19187.

SEQ ID NO: 194 is the determined cDNA sequence for clone #19188.
 SEQ ID NO: 195 is the determined cDNA sequence for clone #19190.
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 5 SEQ ID NO: 198 is the determined cDNA sequence for clone #19193.
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 10 SEQ ID NO: 202 is a first determined cDNA sequence for clone #19200.1.
 SEQ ID NO: 203 is a second determined cDNA sequence for clone
 #19200.2.
 SEQ ID NO: 204 is the determined cDNA sequence for clone #19202.
 SEQ ID NO: 205 is a first determined cDNA sequence for clone #19204.1.
 15 SEQ ID NO: 206 is a second determined cDNA sequence for clone
 #19204.2.
 SEQ ID NO: 207 is the determined cDNA sequence for clone #19205.
 SEQ ID NO: 208 is a first determined cDNA sequence for clone #19206.1.
 SEQ ID NO: 209 is a second determined cDNA sequence for clone
 20 #19206.2.
 SEQ ID NO: 210 is the determined cDNA sequence for clone #19207.
 SEQ ID NO: 211 is the determined cDNA sequence for clone #19208.
 SEQ ID NO: 212 is a first determined cDNA sequence for clone #19211.1.
 SEQ ID NO: 213 is a second determined cDNA sequence for clone
 25 #19211.2.
 SEQ ID NO: 214 is a first determined cDNA sequence for clone #19214.1.
 SEQ ID NO: 215 is a second determined cDNA sequence for clone
 #19214.2.
 SEQ ID NO: 216 is the determined cDNA sequence for clone #19215.

SEQ ID NO: 217 is a first determined cDNA sequence for clone #19217. 2.
 SEQ ID NO: 218 is a second determined cDNA sequence for clone
 #19217.2.

5 SEQ ID NO: 219 is a first determined cDNA sequence for clone #19218.1.
 SEQ ID NO: 220 is a second determined cDNA sequence for clone
 #19218.2.

SEQ ID NO: 221 is a first determined cDNA sequence for clone #19220.1.
 SEQ ID NO: 222 is a second determined cDNA sequence for clone
 #19220.2.

10 SEQ ID NO: 223 is the determined cDNA sequence for clone #22015.
 SEQ ID NO: 224 is the determined cDNA sequence for clone #22017.
 SEQ ID NO: 225 is the determined cDNA sequence for clone #22019.
 SEQ ID NO: 226 is the determined cDNA sequence for clone #22020.
 SEQ ID NO: 227 is the determined cDNA sequence for clone #22023.

15 SEQ ID NO: 228 is the determined cDNA sequence for clone #22026.
 SEQ ID NO: 229 is the determined cDNA sequence for clone #22027.
 SEQ ID NO: 230 is the determined cDNA sequence for clone #22028.
 SEQ ID NO: 231 is the determined cDNA sequence for clone #22032.
 SEQ ID NO: 232 is the determined cDNA sequence for clone #22037.

20 SEQ ID NO: 233 is the determined cDNA sequence for clone #22045.
 SEQ ID NO: 234 is the determined cDNA sequence for clone #22048.
 SEQ ID NO: 235 is the determined cDNA sequence for clone #22050.
 SEQ ID NO: 236 is the determined cDNA sequence for clone #22052.
 SEQ ID NO: 237 is the determined cDNA sequence for clone #22053.

25 SEQ ID NO: 238 is the determined cDNA sequence for clone #22057.
 SEQ ID NO: 239 is the determined cDNA sequence for clone #22066.
 SEQ ID NO: 240 is the determined cDNA sequence for clone #22077.
 SEQ ID NO: 241 is the determined cDNA sequence for clone #22085.
 SEQ ID NO: 242 is the determined cDNA sequence for clone #22105.

SEQ ID NO: 243 is the determined cDNA sequence for clone #22108.
 SEQ ID NO: 244 is the determined cDNA sequence for clone #22109.
 SEQ ID NO: 245 is the determined cDNA sequence for clone #24842.
 SEQ ID NO: 246 is the determined cDNA sequence for clone #24843.
 5 SEQ ID NO: 247 is the determined cDNA sequence for clone #24845.
 SEQ ID NO: 248 is the determined cDNA sequence for clone #24851.
 SEQ ID NO: 249 is the determined cDNA sequence for clone #24852.
 SEQ ID NO: 250 is the determined cDNA sequence for clone #24853.
 SEQ ID NO: 251 is the determined cDNA sequence for clone #24854.
 10 SEQ ID NO: 252 is the determined cDNA sequence for clone #24855.
 SEQ ID NO: 253 is the determined cDNA sequence for clone #24860.
 SEQ ID NO: 254 is the determined cDNA sequence for clone #24864.
 SEQ ID NO: 255 is the determined cDNA sequence for clone #24866.
 SEQ ID NO: 256 is the determined cDNA sequence for clone #24867.
 15 SEQ ID NO: 257 is the determined cDNA sequence for clone #24868.
 SEQ ID NO: 258 is the determined cDNA sequence for clone #24869.
 SEQ ID NO: 259 is the determined cDNA sequence for clone #24870.
 SEQ ID NO: 260 is the determined cDNA sequence for clone #24872.
 SEQ ID NO: 261 is the determined cDNA sequence for clone #24873.
 20 SEQ ID NO: 262 is the determined cDNA sequence for clone #24875.
 SEQ ID NO: 263 is the determined cDNA sequence for clone #24882.
 SEQ ID NO: 264 is the determined cDNA sequence for clone #24885.
 SEQ ID NO: 265 is the determined cDNA sequence for clone #24886.
 SEQ ID NO: 266 is the determined cDNA sequence for clone #24887.
 25 SEQ ID NO: 267 is the determined cDNA sequence for clone #24888.
 SEQ ID NO: 268 is the determined cDNA sequence for clone #24890.
 SEQ ID NO: 269 is the determined cDNA sequence for clone #24896.
 SEQ ID NO: 270 is the determined cDNA sequence for clone #24897.
 SEQ ID NO: 271 is the determined cDNA sequence for clone #24899.

SEQ ID NO: 272 is the determined cDNA sequence for clone #24901.
 SEQ ID NO: 273 is the determined cDNA sequence for clone #24902.
 SEQ ID NO: 274 is the determined cDNA sequence for clone #24906.
 SEQ ID NO: 275 is the determined cDNA sequence for clone #24912.
 5 SEQ ID NO: 276 is the determined cDNA sequence for clone #24913.
 SEQ ID NO: 277 is the determined cDNA sequence for clone #24920.
 SEQ ID NO: 278 is the determined cDNA sequence for clone #24927.
 SEQ ID NO: 279 is the determined cDNA sequence for clone #24930.
 SEQ ID NO: 280 is the determined cDNA sequence for clone #26938.
 10 SEQ ID NO: 281 is the determined cDNA sequence for clone #26939.
 SEQ ID NO: 282 is the determined cDNA sequence for clone #26943.
 SEQ ID NO: 283 is the determined cDNA sequence for clone #26948.
 SEQ ID NO: 284 is the determined cDNA sequence for clone #26951.
 SEQ ID NO: 285 is the determined cDNA sequence for clone #26955.
 15 SEQ ID NO: 286 is the determined cDNA sequence for clone #26956.
 SEQ ID NO: 287 is the determined cDNA sequence for clone #26959.
 SEQ ID NO: 288 is the determined cDNA sequence for clone #26961.
 SEQ ID NO: 289 is the determined cDNA sequence for clone #26962.
 SEQ ID NO: 290 is the determined cDNA sequence for clone #26964.
 20 SEQ ID NO: 291 is the determined cDNA sequence for clone #26966.
 SEQ ID NO: 292 is the determined cDNA sequence for clone #26968.
 SEQ ID NO: 293 is the determined cDNA sequence for clone #26972.
 SEQ ID NO: 294 is the determined cDNA sequence for clone #26973.
 SEQ ID NO: 295 is the determined cDNA sequence for clone #26974.
 25 SEQ ID NO: 296 is the determined cDNA sequence for clone #26976.
 SEQ ID NO: 297 is the determined cDNA sequence for clone #26977.
 SEQ ID NO: 298 is the determined cDNA sequence for clone #26979.
 SEQ ID NO: 299 is the determined cDNA sequence for clone #26980.
 SEQ ID NO: 300 is the determined cDNA sequence for clone #26981.

- SEQ ID NO: 301 is the determined cDNA sequence for clone #26984.
 SEQ ID NO: 302 is the determined cDNA sequence for clone #26985.
 SEQ ID NO: 303 is the determined cDNA sequence for clone #26986.
 SEQ ID NO: 304 is the determined cDNA sequence for clone #26993.
 5 SEQ ID NO: 305 is the determined cDNA sequence for clone #26994.
 SEQ ID NO: 306 is the determined cDNA sequence for clone #26995.
 SEQ ID NO: 307 is the determined cDNA sequence for clone #27003.
 SEQ ID NO: 308 is the determined cDNA sequence for clone #27005.
 SEQ ID NO: 309 is the determined cDNA sequence for clone #27010.
 10 SEQ ID NO: 310 is the determined cDNA sequence for clone #27011.
 SEQ ID NO: 311 is the determined cDNA sequence for clone #27013.
 SEQ ID NO: 312 is the determined cDNA sequence for clone #27016
 SEQ ID NO: 313 is the determined cDNA sequence for clone #27017.
 SEQ ID NO: 314 is the determined cDNA sequence for clone #27019.
 15 SEQ ID NO: 315 is the determined cDNA sequence for clone #27028.
 SEQ ID NO: 316 is the full-length cDNA sequence for clone #19060.
 SEQ ID NO: 317 is the full-length cDNA sequence for clone #18964.
 SEQ ID NO: 318 is the full-length cDNA sequence for clone #18929.
 SEQ ID NO: 319 is the full-length cDNA sequence for clone #18991.
 20 SEQ ID NO: 320 is the full-length cDNA sequence for clone #18996.
 SEQ ID NO: 321 is the full-length cDNA sequence for clone #18966.
 SEQ ID NO: 322 is the full-length cDNA sequence for clone #18951.
 SEQ ID NO: 323 is the full-length cDNA sequence for clone #18973 (also
 known as L516S).
 25 SEQ ID NO: 324 is the amino acid sequence for clone #19060.
 SEQ ID NO: 325 is the amino acid sequence for clone #19063.
 SEQ ID NO: 326 is the amino acid sequence for clone #19077.
 SEQ ID NO: 327 is the amino acid sequence for clone #19110.
 SEQ ID NO: 328 is the amino acid sequence for clone #19122.

5 SEQ ID NO: 329 is the amino acid sequence for clone #19118.
 SEQ ID NO: 330 is the amino acid sequence for clone #19080.
 SEQ ID NO: 331 is the amino acid sequence for clone #19127.
 SEQ ID NO: 332 is the amino acid sequence for clone #19117.
 SEQ ID NO: 333 is the amino acid sequence for clone #19095, also referred
 to L549S.

10 SEQ ID NO: 334 is the amino acid sequence for clone #18964.
 SEQ ID NO: 335 is the amino acid sequence for clone #18929.
 SEQ ID NO: 336 is the amino acid sequence for clone #18991.
 SEQ ID NO: 337 is the amino acid sequence for clone #18996.
 SEQ ID NO: 338 is the amino acid sequence for clone #18966.
 SEQ ID NO: 339 is the amino acid sequence for clone #18951.
 SEQ ID NO: 340 is the amino acid sequence for clone #18973.
 SEQ ID NO: 341 is the determined cDNA sequence for clone 26461.
 15 SEQ ID NO: 342 is the determined cDNA sequence for clone 26462.
 SEQ ID NO: 343 is the determined cDNA sequence for clone 26463.
 SEQ ID NO: 344 is the determined cDNA sequence for clone 26464.
 SEQ ID NO: 345 is the determined cDNA sequence for clone 26465.
 SEQ ID NO: 346 is the determined cDNA sequence for clone 26466.
 20 SEQ ID NO: 347 is the determined cDNA sequence for clone 26467.
 SEQ ID NO: 348 is the determined cDNA sequence for clone 26468.
 SEQ ID NO: 349 is the determined cDNA sequence for clone 26469.
 SEQ ID NO: 350 is the determined cDNA sequence for clone 26470.
 SEQ ID NO: 351 is the determined cDNA sequence for clone 26471.
 25 SEQ ID NO: 352 is the determined cDNA sequence for clone 26472.
 SEQ ID NO: 353 is the determined cDNA sequence for clone 26474.
 SEQ ID NO: 354 is the determined cDNA sequence for clone 26475.
 SEQ ID NO: 355 is the determined cDNA sequence for clone 26476.
 SEQ ID NO: 356 is the determined cDNA sequence for clone 26477.

SEQ ID NO: 357 is the determined cDNA sequence for clone 26478.
 SEQ ID NO: 358 is the determined cDNA sequence for clone 26479.
 SEQ ID NO: 359 is the determined cDNA sequence for clone 26480.
 SEQ ID NO: 360 is the determined cDNA sequence for clone 26481.
 5 SEQ ID NO: 361 is the determined cDNA sequence for clone 26482
 SEQ ID NO: 362 is the determined cDNA sequence for clone 26483.
 SEQ ID NO: 363 is the determined cDNA sequence for clone 26484.
 SEQ ID NO: 364 is the determined cDNA sequence for clone 26485.
 SEQ ID NO: 365 is the determined cDNA sequence for clone 26486.
 10 SEQ ID NO: 366 is the determined cDNA sequence for clone 26487.
 SEQ ID NO: 367 is the determined cDNA sequence for clone 26488.
 SEQ ID NO: 368 is the determined cDNA sequence for clone 26489.
 SEQ ID NO: 369 is the determined cDNA sequence for clone 26490.
 SEQ ID NO: 370 is the determined cDNA sequence for clone 26491.
 15 SEQ ID NO: 371 is the determined cDNA sequence for clone 26492.
 SEQ ID NO: 372 is the determined cDNA sequence for clone 26493.
 SEQ ID NO: 373 is the determined cDNA sequence for clone 26494.
 SEQ ID NO: 374 is the determined cDNA sequence for clone 26495.
 SEQ ID NO: 375 is the determined cDNA sequence for clone 26496.
 20 SEQ ID NO: 376 is the determined cDNA sequence for clone 26497.
 SEQ ID NO: 377 is the determined cDNA sequence for clone 26498.
 SEQ ID NO: 378 is the determined cDNA sequence for clone 26499.
 SEQ ID NO: 379 is the determined cDNA sequence for clone 26500.
 SEQ ID NO: 380 is the determined cDNA sequence for clone 26501.
 25 SEQ ID NO: 381 is the determined cDNA sequence for clone 26502.
 SEQ ID NO: 382 is the determined cDNA sequence for clone 26503.
 SEQ ID NO: 383 is the determined cDNA sequence for clone 26504.
 SEQ ID NO: 384 is the determined cDNA sequence for clone 26505.
 SEQ ID NO: 385 is the determined cDNA sequence for clone 26506.

SEQ ID NO: 386 is the determined cDNA sequence for clone 26507.
 SEQ ID NO: 387 is the determined cDNA sequence for clone 26508.
 SEQ ID NO: 388 is the determined cDNA sequence for clone 26509.
 SEQ ID NO: 389 is the determined cDNA sequence for clone 26511.
 5 SEQ ID NO: 390 is the determined cDNA sequence for clone 26513.
 SEQ ID NO: 391 is the determined cDNA sequence for clone 26514.
 SEQ ID NO: 392 is the determined cDNA sequence for clone 26515.
 SEQ ID NO: 393 is the determined cDNA sequence for clone 26516.
 SEQ ID NO: 394 is the determined cDNA sequence for clone 26517.
 10 SEQ ID NO: 395 is the determined cDNA sequence for clone 26518.
 SEQ ID NO: 396 is the determined cDNA sequence for clone 26519.
 SEQ ID NO: 397 is the determined cDNA sequence for clone 26520.
 SEQ ID NO: 398 is the determined cDNA sequence for clone 26521.
 SEQ ID NO: 399 is the determined cDNA sequence for clone 26522.
 15 SEQ ID NO: 400 is the determined cDNA sequence for clone 26523.
 SEQ ID NO: 401 is the determined cDNA sequence for clone 26524.
 SEQ ID NO: 402 is the determined cDNA sequence for clone 26526.
 SEQ ID NO: 403 is the determined cDNA sequence for clone 26527.
 SEQ ID NO: 404 is the determined cDNA sequence for clone 26528.
 20 SEQ ID NO: 405 is the determined cDNA sequence for clone 26529.
 SEQ ID NO: 406 is the determined cDNA sequence for clone 26530.
 SEQ ID NO: 407 is the determined cDNA sequence for clone 26532.
 SEQ ID NO: 408 is the determined cDNA sequence for clone 26533.
 SEQ ID NO: 409 is the determined cDNA sequence for clone 26534.
 25 SEQ ID NO: 410 is the determined cDNA sequence for clone 26535.
 SEQ ID NO: 411 is the determined cDNA sequence for clone 26536.
 SEQ ID NO: 412 is the determined cDNA sequence for clone 26537.
 SEQ ID NO: 413 is the determined cDNA sequence for clone 26538.
 SEQ ID NO: 414 is the determined cDNA sequence for clone 26540.

SEQ ID NO: 415 is the determined cDNA sequence for clone 26541.
 SEQ ID NO: 416 is the determined cDNA sequence for clone 26542.
 SEQ ID NO: 417 is the determined cDNA sequence for clone 26543.
 SEQ ID NO: 418 is the determined cDNA sequence for clone 26544.
 5 SEQ ID NO: 419 is the determined cDNA sequence for clone 26546.
 SEQ ID NO: 420 is the determined cDNA sequence for clone 26547.
 SEQ ID NO: 421 is the determined cDNA sequence for clone 26548.
 SEQ ID NO: 422 is the determined cDNA sequence for clone 26549.
 SEQ ID NO: 423 is the determined cDNA sequence for clone 26550.
 10 SEQ ID NO: 424 is the determined cDNA sequence for clone 26551.
 SEQ ID NO: 425 is the determined cDNA sequence for clone 26552.
 SEQ ID NO: 426 is the determined cDNA sequence for clone 26553.
 SEQ ID NO: 427 is the determined cDNA sequence for clone 26554.
 SEQ ID NO: 428 is the determined cDNA sequence for clone 26556.
 15 SEQ ID NO: 429 is the determined cDNA sequence for clone 26557.
 SEQ ID NO: 430 is the determined cDNA sequence for clone 27631.
 SEQ ID NO: 431 is the determined cDNA sequence for clone 27632.
 SEQ ID NO: 432 is the determined cDNA sequence for clone 27633.
 SEQ ID NO: 433 is the determined cDNA sequence for clone 27635.
 20 SEQ ID NO: 434 is the determined cDNA sequence for clone 27636.
 SEQ ID NO: 435 is the determined cDNA sequence for clone 27637.
 SEQ ID NO: 436 is the determined cDNA sequence for clone 27638.
 SEQ ID NO: 437 is the determined cDNA sequence for clone 27639.
 SEQ ID NO: 438 is the determined cDNA sequence for clone 27640.
 25 SEQ ID NO: 439 is the determined cDNA sequence for clone 27641.
 SEQ ID NO: 440 is the determined cDNA sequence for clone 27642.
 SEQ ID NO: 441 is the determined cDNA sequence for clone 27644.
 SEQ ID NO: 442 is the determined cDNA sequence for clone 27646.
 SEQ ID NO: 443 is the determined cDNA sequence for clone 27647.

SEQ ID NO: 444 is the determined cDNA sequence for clone 27649.
 SEQ ID NO: 445 is the determined cDNA sequence for clone 27650.
 SEQ ID NO: 446 is the determined cDNA sequence for clone 27651.
 SEQ ID NO: 447 is the determined cDNA sequence for clone 27652.
 5 SEQ ID NO: 448 is the determined cDNA sequence for clone 27654.
 SEQ ID NO: 449 is the determined cDNA sequence for clone 27655.
 SEQ ID NO: 450 is the determined cDNA sequence for clone 27657.
 SEQ ID NO: 451 is the determined cDNA sequence for clone 27659.
 SEQ ID NO: 452 is the determined cDNA sequence for clone 27665.
 10 SEQ ID NO: 453 is the determined cDNA sequence for clone 27666.
 SEQ ID NO: 454 is the determined cDNA sequence for clone 27668.
 SEQ ID NO: 455 is the determined cDNA sequence for clone 27670.
 SEQ ID NO: 456 is the determined cDNA sequence for clone 27671.
 SEQ ID NO: 457 is the determined cDNA sequence for clone 27672.
 15 SEQ ID NO: 458 is the determined cDNA sequence for clone 27674.
 SEQ ID NO: 459 is the determined cDNA sequence for clone 27677.
 SEQ ID NO: 460 is the determined cDNA sequence for clone 27681.
 SEQ ID NO: 461 is the determined cDNA sequence for clone 27682.
 SEQ ID NO: 462 is the determined cDNA sequence for clone 27683.
 20 SEQ ID NO: 463 is the determined cDNA sequence for clone 27686.
 SEQ ID NO: 464 is the determined cDNA sequence for clone 27688.
 SEQ ID NO: 465 is the determined cDNA sequence for clone 27689.
 SEQ ID NO: 466 is the determined cDNA sequence for clone 27690.
 SEQ ID NO: 467 is the determined cDNA sequence for clone 27693.
 25 SEQ ID NO: 468 is the determined cDNA sequence for clone 27699.
 SEQ ID NO: 469 is the determined cDNA sequence for clone 27700.
 SEQ ID NO: 470 is the determined cDNA sequence for clone 27702.
 SEQ ID NO: 471 is the determined cDNA sequence for clone 27705.
 SEQ ID NO: 472 is the determined cDNA sequence for clone 27706.

SEQ ID NO: 473 is the determined cDNA sequence for clone 27707.
 SEQ ID NO: 474 is the determined cDNA sequence for clone 27708.
 SEQ ID NO: 475 is the determined cDNA sequence for clone 27709.
 SEQ ID NO: 476 is the determined cDNA sequence for clone 27710.
 5 SEQ ID NO: 477 is the determined cDNA sequence for clone 27711.
 SEQ ID NO: 478 is the determined cDNA sequence for clone 27712.
 SEQ ID NO: 479 is the determined cDNA sequence for clone 27713.
 SEQ ID NO: 480 is the determined cDNA sequence for clone 27714.
 SEQ ID NO: 481 is the determined cDNA sequence for clone 27715.
 10 SEQ ID NO: 482 is the determined cDNA sequence for clone 27716.
 SEQ ID NO: 483 is the determined cDNA sequence for clone 27717.
 SEQ ID NO: 484 is the determined cDNA sequence for clone 27718.
 SEQ ID NO: 485 is the determined cDNA sequence for clone 27719.
 SEQ ID NO: 486 is the determined cDNA sequence for clone 27720.
 15 SEQ ID NO: 487 is the determined cDNA sequence for clone 27722.
 SEQ ID NO: 488 is the determined cDNA sequence for clone 27723.
 SEQ ID NO: 489 is the determined cDNA sequence for clone 27724.
 SEQ ID NO: 490 is the determined cDNA sequence for clone 27726.
 SEQ ID NO: 491 is the determined cDNA sequence for clone 25015.
 20 SEQ ID NO: 492 is the determined cDNA sequence for clone 25016.
 SEQ ID NO: 493 is the determined cDNA sequence for clone 25017.
 SEQ ID NO: 494 is the determined cDNA sequence for clone 25018.
 SEQ ID NO: 495 is the determined cDNA sequence for clone 25030.
 SEQ ID NO: 496 is the determined cDNA sequence for clone 25033.
 25 SEQ ID NO: 497 is the determined cDNA sequence for clone 25034.
 SEQ ID NO: 498 is the determined cDNA sequence for clone 25035.
 SEQ ID NO: 499 is the determined cDNA sequence for clone 25036.
 SEQ ID NO: 500 is the determined cDNA sequence for clone 25037.
 SEQ ID NO: 501 is the determined cDNA sequence for clone 25038.

SEQ ID NO: 502 is the determined cDNA sequence for clone 25039.
 SEQ ID NO: 503 is the determined cDNA sequence for clone 25040.
 SEQ ID NO: 504 is the determined cDNA sequence for clone 25042.
 SEQ ID NO: 505 is the determined cDNA sequence for clone 25043.
 5 SEQ ID NO: 506 is the determined cDNA sequence for clone 25044.
 SEQ ID NO: 507 is the determined cDNA sequence for clone 25045.
 SEQ ID NO: 508 is the determined cDNA sequence for clone 25047.
 SEQ ID NO: 509 is the determined cDNA sequence for clone 25048.
 SEQ ID NO: 510 is the determined cDNA sequence for clone 25049.
 10 SEQ ID NO: 511 is the determined cDNA sequence for clone 25185.
 SEQ ID NO: 512 is the determined cDNA sequence for clone 25186.
 SEQ ID NO: 513 is the determined cDNA sequence for clone 25187.
 SEQ ID NO: 514 is the determined cDNA sequence for clone 25188.
 SEQ ID NO: 515 is the determined cDNA sequence for clone 25189.
 15 SEQ ID NO: 516 is the determined cDNA sequence for clone 25190.
 SEQ ID NO: 517 is the determined cDNA sequence for clone 25193.
 SEQ ID NO: 518 is the determined cDNA sequence for clone 25194.
 SEQ ID NO: 519 is the determined cDNA sequence for clone 25196.
 SEQ ID NO: 520 is the determined cDNA sequence for clone 25198.
 20 SEQ ID NO: 521 is the determined cDNA sequence for clone 25199.
 SEQ ID NO: 522 is the determined cDNA sequence for clone 25200.
 SEQ ID NO: 523 is the determined cDNA sequence for clone 25202.
 SEQ ID NO: 524 is the determined cDNA sequence for clone 25364.
 SEQ ID NO: 525 is the determined cDNA sequence for clone 25366.
 25 SEQ ID NO: 526 is the determined cDNA sequence for clone 25367.
 SEQ ID NO: 527 is the determined cDNA sequence for clone 25368.
 SEQ ID NO: 528 is the determined cDNA sequence for clone 25369.
 SEQ ID NO: 529 is the determined cDNA sequence for clone 25370.
 SEQ ID NO: 530 is the determined cDNA sequence for clone 25371.

SEQ ID NO: 531 is the determined cDNA sequence for clone 25372.
 SEQ ID NO: 532 is the determined cDNA sequence for clone 25373.
 SEQ ID NO: 533 is the determined cDNA sequence for clone 25374.
 SEQ ID NO: 534 is the determined cDNA sequence for clone 25376.
 5 SEQ ID NO: 535 is the determined cDNA sequence for clone 25377.
 SEQ ID NO: 536 is the determined cDNA sequence for clone 25378.
 SEQ ID NO: 537 is the determined cDNA sequence for clone 25379.
 SEQ ID NO: 538 is the determined cDNA sequence for clone 25380.
 SEQ ID NO: 539 is the determined cDNA sequence for clone 25381.
 10 SEQ ID NO: 540 is the determined cDNA sequence for clone 25382.
 SEQ ID NO: 541 is the determined cDNA sequence for clone 25383.
 SEQ ID NO: 542 is the determined cDNA sequence for clone 25385.
 SEQ ID NO: 543 is the determined cDNA sequence for clone 25386.
 SEQ ID NO: 544 is the determined cDNA sequence for clone 25387.
 15 SEQ ID NO: 545 is the determined cDNA sequence for clone 26013.
 SEQ ID NO: 546 is the determined cDNA sequence for clone 26014.
 SEQ ID NO: 547 is the determined cDNA sequence for clone 26016.
 SEQ ID NO: 548 is the determined cDNA sequence for clone 26017.
 SEQ ID NO: 549 is the determined cDNA sequence for clone 26018.
 20 SEQ ID NO: 550 is the determined cDNA sequence for clone 26019.
 SEQ ID NO: 551 is the determined cDNA sequence for clone 26020.
 SEQ ID NO: 552 is the determined cDNA sequence for clone 26021.
 SEQ ID NO: 553 is the determined cDNA sequence for clone 26022.
 SEQ ID NO: 554 is the determined cDNA sequence for clone 26027.
 25 SEQ ID NO: 555 is the determined cDNA sequence for clone 26197.
 SEQ ID NO: 556 is the determined cDNA sequence for clone 26199.
 SEQ ID NO: 557 is the determined cDNA sequence for clone 26201.
 SEQ ID NO: 558 is the determined cDNA sequence for clone 26202.
 SEQ ID NO: 559 is the determined cDNA sequence for clone 26203.

SEQ ID NO: 560 is the determined cDNA sequence for clone 26204.
 SEQ ID NO: 561 is the determined cDNA sequence for clone 26205.
 SEQ ID NO: 562 is the determined cDNA sequence for clone 26206.
 SEQ ID NO: 563 is the determined cDNA sequence for clone 26208.
 5 SEQ ID NO: 564 is the determined cDNA sequence for clone 26211.
 SEQ ID NO: 565 is the determined cDNA sequence for clone 26212.
 SEQ ID NO: 566 is the determined cDNA sequence for clone 26213.
 SEQ ID NO: 567 is the determined cDNA sequence for clone 26214.
 SEQ ID NO: 568 is the determined cDNA sequence for clone 26215.
 10 SEQ ID NO: 569 is the determined cDNA sequence for clone 26216.
 SEQ ID NO: 570 is the determined cDNA sequence for clone 26217.
 SEQ ID NO: 571 is the determined cDNA sequence for clone 26218.
 SEQ ID NO: 572 is the determined cDNA sequence for clone 26219.
 SEQ ID NO: 573 is the determined cDNA sequence for clone 26220.
 15 SEQ ID NO: 574 is the determined cDNA sequence for clone 26221.
 SEQ ID NO: 575 is the determined cDNA sequence for clone 26224.
 SEQ ID NO: 576 is the determined cDNA sequence for clone 26225.
 SEQ ID NO: 577 is the determined cDNA sequence for clone 26226.
 SEQ ID NO: 578 is the determined cDNA sequence for clone 26227.
 20 SEQ ID NO: 579 is the determined cDNA sequence for clone 26228.
 SEQ ID NO: 580 is the determined cDNA sequence for clone 26230.
 SEQ ID NO: 581 is the determined cDNA sequence for clone 26231.
 SEQ ID NO: 582 is the determined cDNA sequence for clone 26234.
 SEQ ID NO: 583 is the determined cDNA sequence for clone 26236.
 25 SEQ ID NO: 584 is the determined cDNA sequence for clone 26237.
 SEQ ID NO: 585 is the determined cDNA sequence for clone 26239.
 SEQ ID NO: 586 is the determined cDNA sequence for clone 26240.
 SEQ ID NO: 587 is the determined cDNA sequence for clone 26241.
 SEQ ID NO: 588 is the determined cDNA sequence for clone 26242.

SEQ ID NO: 589 is the determined cDNA sequence for clone 26246.
 SEQ ID NO: 590 is the determined cDNA sequence for clone 26247.
 SEQ ID NO: 591 is the determined cDNA sequence for clone 26248.
 SEQ ID NO: 592 is the determined cDNA sequence for clone 26249.
 5 SEQ ID NO: 593 is the determined cDNA sequence for clone 26250.
 SEQ ID NO: 594 is the determined cDNA sequence for clone 26251.
 SEQ ID NO: 595 is the determined cDNA sequence for clone 26252.
 SEQ ID NO: 596 is the determined cDNA sequence for clone 26253.
 SEQ ID NO: 597 is the determined cDNA sequence for clone 26254.
 10 SEQ ID NO: 598 is the determined cDNA sequence for clone 26255.
 SEQ ID NO: 599 is the determined cDNA sequence for clone 26256.
 SEQ ID NO: 600 is the determined cDNA sequence for clone 26257.
 SEQ ID NO: 601 is the determined cDNA sequence for clone 26259.
 SEQ ID NO: 602 is the determined cDNA sequence for clone 26260.
 15 SEQ ID NO: 603 is the determined cDNA sequence for clone 26261.
 SEQ ID NO: 604 is the determined cDNA sequence for clone 26262.
 SEQ ID NO: 605 is the determined cDNA sequence for clone 26263.
 SEQ ID NO: 606 is the determined cDNA sequence for clone 26264.
 SEQ ID NO: 607 is the determined cDNA sequence for clone 26265.
 20 SEQ ID NO: 608 is the determined cDNA sequence for clone 26266.
 SEQ ID NO: 609 is the determined cDNA sequence for clone 26268.
 SEQ ID NO: 610 is the determined cDNA sequence for clone 26269.
 SEQ ID NO: 611 is the determined cDNA sequence for clone 26271.
 SEQ ID NO: 612 is the determined cDNA sequence for clone 26273.
 25 SEQ ID NO: 613 is the determined cDNA sequence for clone 26810.
 SEQ ID NO: 614 is the determined cDNA sequence for clone 26811.
 SEQ ID NO: 615 is the determined cDNA sequence for clone 26812.1.
 SEQ ID NO: 616 is the determined cDNA sequence for clone 26812.2.
 SEQ ID NO: 617 is the determined cDNA sequence for clone 26813.

SEQ ID NO: 618 is the determined cDNA sequence for clone 26814.
 SEQ ID NO: 619 is the determined cDNA sequence for clone 26815.
 SEQ ID NO: 620 is the determined cDNA sequence for clone 26816.
 SEQ ID NO: 621 is the determined cDNA sequence for clone 26818.
 5 SEQ ID NO: 622 is the determined cDNA sequence for clone 26819.
 SEQ ID NO: 623 is the determined cDNA sequence for clone 26820.
 SEQ ID NO: 624 is the determined cDNA sequence for clone 26821.
 SEQ ID NO: 625 is the determined cDNA sequence for clone 26822.
 SEQ ID NO: 626 is the determined cDNA sequence for clone 26824.
 10 SEQ ID NO: 627 is the determined cDNA sequence for clone 26825.
 SEQ ID NO: 628 is the determined cDNA sequence for clone 26826.
 SEQ ID NO: 629 is the determined cDNA sequence for clone 26827.
 SEQ ID NO: 630 is the determined cDNA sequence for clone 26829.
 SEQ ID NO: 631 is the determined cDNA sequence for clone 26830.
 15 SEQ ID NO: 632 is the determined cDNA sequence for clone 26831.
 SEQ ID NO: 633 is the determined cDNA sequence for clone 26832.
 SEQ ID NO: 634 is the determined cDNA sequence for clone 26835.
 SEQ ID NO: 635 is the determined cDNA sequence for clone 26836.
 SEQ ID NO: 636 is the determined cDNA sequence for clone 26837.
 20 SEQ ID NO: 637 is the determined cDNA sequence for clone 26839.
 SEQ ID NO: 638 is the determined cDNA sequence for clone 26841.
 SEQ ID NO: 639 is the determined cDNA sequence for clone 26843.
 SEQ ID NO: 640 is the determined cDNA sequence for clone 26844.
 SEQ ID NO: 641 is the determined cDNA sequence for clone 26845.
 25 SEQ ID NO: 642 is the determined cDNA sequence for clone 26846.
 SEQ ID NO: 643 is the determined cDNA sequence for clone 26847.
 SEQ ID NO: 644 is the determined cDNA sequence for clone 26848.
 SEQ ID NO: 645 is the determined cDNA sequence for clone 26849.
 SEQ ID NO: 646 is the determined cDNA sequence for clone 26850.

SEQ ID NO: 647 is the determined cDNA sequence for clone 26851.
 SEQ ID NO: 648 is the determined cDNA sequence for clone 26852.
 SEQ ID NO: 649 is the determined cDNA sequence for clone 26853.
 SEQ ID NO: 650 is the determined cDNA sequence for clone 26854.
 5 SEQ ID NO: 651 is the determined cDNA sequence for clone 26856.
 SEQ ID NO: 652 is the determined cDNA sequence for clone 26857.
 SEQ ID NO: 653 is the determined cDNA sequence for clone 26858.
 SEQ ID NO: 654 is the determined cDNA sequence for clone 26859.
 SEQ ID NO: 655 is the determined cDNA sequence for clone 26860.
 10 SEQ ID NO: 656 is the determined cDNA sequence for clone 26862.
 SEQ ID NO: 657 is the determined cDNA sequence for clone 26863.
 SEQ ID NO: 658 is the determined cDNA sequence for clone 26864.
 SEQ ID NO: 659 is the determined cDNA sequence for clone 26865.
 SEQ ID NO: 660 is the determined cDNA sequence for clone 26867.
 15 SEQ ID NO: 661 is the determined cDNA sequence for clone 26868.
 SEQ ID NO: 662 is the determined cDNA sequence for clone 26871.
 SEQ ID NO: 663 is the determined cDNA sequence for clone 26873.
 SEQ ID NO: 664 is the determined cDNA sequence for clone 26875.
 SEQ ID NO: 665 is the determined cDNA sequence for clone 26876.
 20 SEQ ID NO: 666 is the determined cDNA sequence for clone 26877.
 SEQ ID NO: 667 is the determined cDNA sequence for clone 26878.
 SEQ ID NO: 668 is the determined cDNA sequence for clone 26880.
 SEQ ID NO: 669 is the determined cDNA sequence for clone 26882.
 SEQ ID NO: 670 is the determined cDNA sequence for clone 26883.
 25 SEQ ID NO: 671 is the determined cDNA sequence for clone 26884.
 SEQ ID NO: 672 is the determined cDNA sequence for clone 26885.
 SEQ ID NO: 673 is the determined cDNA sequence for clone 26886.
 SEQ ID NO: 674 is the determined cDNA sequence for clone 26887.
 SEQ ID NO: 675 is the determined cDNA sequence for clone 26888.

SEQ ID NO: 676 is the determined cDNA sequence for clone 26889.
 SEQ ID NO: 677 is the determined cDNA sequence for clone 26890.
 SEQ ID NO: 678 is the determined cDNA sequence for clone 26892.
 SEQ ID NO: 679 is the determined cDNA sequence for clone 26894.
 5 SEQ ID NO: 680 is the determined cDNA sequence for clone 26895.
 SEQ ID NO: 681 is the determined cDNA sequence for clone 26897.
 SEQ ID NO: 682 is the determined cDNA sequence for clone 26898.
 SEQ ID NO: 683 is the determined cDNA sequence for clone 26899.
 SEQ ID NO: 684 is the determined cDNA sequence for clone 26900.
 10 SEQ ID NO: 685 is the determined cDNA sequence for clone 26901.
 SEQ ID NO: 686 is the determined cDNA sequence for clone 26903.
 SEQ ID NO: 687 is the determined cDNA sequence for clone 26905.
 SEQ ID NO: 688 is the determined cDNA sequence for clone 26906.
 SEQ ID NO: 689 is the determined cDNA sequence for clone 26708.
 15 SEQ ID NO: 690 is the determined cDNA sequence for clone 26709.
 SEQ ID NO: 691 is the determined cDNA sequence for clone 26710.
 SEQ ID NO: 692 is the determined cDNA sequence for clone 26711.
 SEQ ID NO: 693 is the determined cDNA sequence for clone 26712.
 SEQ ID NO: 694 is the determined cDNA sequence for clone 26713.
 20 SEQ ID NO: 695 is the determined cDNA sequence for clone 26714.
 SEQ ID NO: 696 is the determined cDNA sequence for clone 26715.
 SEQ ID NO: 697 is the determined cDNA sequence for clone 26716.
 SEQ ID NO: 698 is the determined cDNA sequence for clone 26717.
 SEQ ID NO: 699 is the determined cDNA sequence for clone 26718.
 25 SEQ ID NO: 700 is the determined cDNA sequence for clone 26719.
 SEQ ID NO: 701 is the determined cDNA sequence for clone 26720.
 SEQ ID NO: 702 is the determined cDNA sequence for clone 26721.
 SEQ ID NO: 703 is the determined cDNA sequence for clone 26722.
 SEQ ID NO: 704 is the determined cDNA sequence for clone 26723.

SEQ ID NO: 705 is the determined cDNA sequence for clone 26724.
 SEQ ID NO: 706 is the determined cDNA sequence for clone 26725.
 SEQ ID NO: 707 is the determined cDNA sequence for clone 26726.
 SEQ ID NO: 708 is the determined cDNA sequence for clone 26727.
 5 SEQ ID NO: 709 is the determined cDNA sequence for clone 26728.
 SEQ ID NO: 710 is the determined cDNA sequence for clone 26729.
 SEQ ID NO: 711 is the determined cDNA sequence for clone 26730.
 SEQ ID NO: 712 is the determined cDNA sequence for clone 26731.
 SEQ ID NO: 713 is the determined cDNA sequence for clone 26732.
 10 SEQ ID NO: 714 is the determined cDNA sequence for clone 26733.1.
 SEQ ID NO: 715 is the determined cDNA sequence for clone 26733.2.
 SEQ ID NO: 716 is the determined cDNA sequence for clone 26734.
 SEQ ID NO: 717 is the determined cDNA sequence for clone 26735.
 SEQ ID NO: 718 is the determined cDNA sequence for clone 26736.
 15 SEQ ID NO: 719 is the determined cDNA sequence for clone 26737.
 SEQ ID NO: 720 is the determined cDNA sequence for clone 26738.
 SEQ ID NO: 721 is the determined cDNA sequence for clone 26739.
 SEQ ID NO: 722 is the determined cDNA sequence for clone 26741.
 SEQ ID NO: 723 is the determined cDNA sequence for clone 26742.
 20 SEQ ID NO: 724 is the determined cDNA sequence for clone 26743.
 SEQ ID NO: 725 is the determined cDNA sequence for clone 26744.
 SEQ ID NO: 726 is the determined cDNA sequence for clone 26745.
 SEQ ID NO: 727 is the determined cDNA sequence for clone 26746.
 SEQ ID NO: 728 is the determined cDNA sequence for clone 26747.
 25 SEQ ID NO: 729 is the determined cDNA sequence for clone 26748.
 SEQ ID NO: 730 is the determined cDNA sequence for clone 26749.
 SEQ ID NO: 731 is the determined cDNA sequence for clone 26750.
 SEQ ID NO: 732 is the determined cDNA sequence for clone 26751.
 SEQ ID NO: 733 is the determined cDNA sequence for clone 26752.

SEQ ID NO: 734 is the determined cDNA sequence for clone 26753.
 SEQ ID NO: 735 is the determined cDNA sequence for clone 26754.
 SEQ ID NO: 736 is the determined cDNA sequence for clone 26755.
 SEQ ID NO: 737 is the determined cDNA sequence for clone 26756.
 5 SEQ ID NO: 738 is the determined cDNA sequence for clone 26757.
 SEQ ID NO: 739 is the determined cDNA sequence for clone 26758.
 SEQ ID NO: 740 is the determined cDNA sequence for clone 26759.
 SEQ ID NO: 741 is the determined cDNA sequence for clone 26760.
 SEQ ID NO: 742 is the determined cDNA sequence for clone 26761.
 10 SEQ ID NO: 743 is the determined cDNA sequence for clone 26762.
 SEQ ID NO: 744 is the determined cDNA sequence for clone 26763.
 SEQ ID NO: 745 is the determined cDNA sequence for clone 26764.
 SEQ ID NO: 746 is the determined cDNA sequence for clone 26765.
 SEQ ID NO: 747 is the determined cDNA sequence for clone 26766.
 15 SEQ ID NO: 748 is the determined cDNA sequence for clone 26767.
 SEQ ID NO: 749 is the determined cDNA sequence for clone 26768.
 SEQ ID NO: 750 is the determined cDNA sequence for clone 26769.
 SEQ ID NO: 751 is the determined cDNA sequence for clone 26770.
 SEQ ID NO: 752 is the determined cDNA sequence for clone 26771.
 20 SEQ ID NO: 753 is the determined cDNA sequence for clone 26772.
 SEQ ID NO: 754 is the determined cDNA sequence for clone 26773.
 SEQ ID NO: 755 is the determined cDNA sequence for clone 26774.
 SEQ ID NO: 756 is the determined cDNA sequence for clone 26775.
 SEQ ID NO: 757 is the determined cDNA sequence for clone 26776.
 25 SEQ ID NO: 758 is the determined cDNA sequence for clone 26777.
 SEQ ID NO: 759 is the determined cDNA sequence for clone 26778.
 SEQ ID NO: 760 is the determined cDNA sequence for clone 26779.
 SEQ ID NO: 761 is the determined cDNA sequence for clone 26781.
 SEQ ID NO: 762 is the determined cDNA sequence for clone 26782.

SEQ ID NO: 763 is the determined cDNA sequence for clone 26783.
 SEQ ID NO: 764 is the determined cDNA sequence for clone 26784.
 SEQ ID NO: 765 is the determined cDNA sequence for clone 26785.
 SEQ ID NO: 766 is the determined cDNA sequence for clone 26786.
 5 SEQ ID NO: 767 is the determined cDNA sequence for clone 26787.
 SEQ ID NO: 768 is the determined cDNA sequence for clone 26788.
 SEQ ID NO: 769 is the determined cDNA sequence for clone 26790.
 SEQ ID NO: 770 is the determined cDNA sequence for clone 26791.
 SEQ ID NO: 771 is the determined cDNA sequence for clone 26792.
 10 SEQ ID NO: 772 is the determined cDNA sequence for clone 26793.
 SEQ ID NO: 773 is the determined cDNA sequence for clone 26794.
 SEQ ID NO: 774 is the determined cDNA sequence for clone 26795.
 SEQ ID NO: 775 is the determined cDNA sequence for clone 26796.
 SEQ ID NO: 776 is the determined cDNA sequence for clone 26797.
 15 SEQ ID NO: 777 is the determined cDNA sequence for clone 26798.
 SEQ ID NO: 778 is the determined cDNA sequence for clone 26800.
 SEQ ID NO: 779 is the determined cDNA sequence for clone 26801.
 SEQ ID NO: 780 is the determined cDNA sequence for clone 26802.
 SEQ ID NO: 781 is the determined cDNA sequence for clone 26803.
 20 SEQ ID NO: 782 is the determined cDNA sequence for clone 26804.
 SEQ ID NO: 783 is the amino acid sequence for L773P.
 SEQ ID NO: 784 is the determined DNA sequence of the L773P expression
 construct.
 SEQ ID NO: 785 is the determined DNA sequence of the L773PA
 25 expression construct.
 SEQ ID NO: 786 is a predicted amino acid sequence for L552S.
 SEQ ID NO: 787 is a predicted amino acid sequence for L840P.
 SEQ ID NO: 788 is the full-length cDNA sequence for L548S.
 SEQ ID NO: 789 is the amino acid sequence encoded by SEQ ID NO: 788.

SEQ ID NO: 790 is an extended cDNA sequence for L552S.

SEQ ID NO: 791 is the predicted amino acid sequence encoded by the cDNA sequence of SEQ ID NO: 790.

5 SEQ ID NO: 792 is the determined cDNA sequence for an isoform of L552S.

SEQ ID NO: 793 is the predicted amino acid sequence encoded by SEQ ID NO: 792.

SEQ ID NO: 794 is an extended cDNA sequence for L840P.

10 SEQ ID NO: 795 is the predicted amino acid sequence encoded by SEQ ID NO: 794.

SEQ ID NO: 796 is an extended cDNA sequence for L801P.

SEQ ID NO: 797 is a first predicted amino acid sequence encoded by SEQ ID NO: 796.

15 SEQ ID NO: 798 is a second predicted amino acid sequence encoded by SEQ ID NO: 796.

SEQ ID NO: 799 is a third predicted amino acid sequence encoded by SEQ ID NO: 796.

SEQ ID NO: 800 is the determined full-length sequence for L844P.

SEQ ID NO: 801 is the 5' consensus cDNA sequence for L551S.

20 SEQ ID NO: 802 is the 3' consensus cDNA sequence for L551S.

SEQ ID NO: 803 is the cDNA sequence for STY8.

SEQ ID NO: 804 is an extended cDNA sequence for L551S.

SEQ ID NO: 805 is the amino acid sequence for STY8.

SEQ ID NO: 806 is the extended amino acid sequence for L551S.

25 SEQ ID NO: 807 is the determined full-length cDNA sequence for L773P.

SEQ ID NO: 808 is the full-length cDNA sequence of L552S.

SEQ ID NO: 809 is the full-length amino acid sequence of L552S.

SEQ ID NO: 810 is the determined cDNA sequence of clone 50989.

SEQ ID NO: 811 is the determined cDNA sequence of clone 50990.

SEQ ID NO: 812 is the determined cDNA sequence of clone 50992.

SEQ ID NO: 813-824 are the determined cDNA sequences for clones isolated from lung tumor tissue.

SEQ ID NO: 825 is the determined cDNA sequence for the full-length
5 L551S clone 54305.

SEQ ID NO: 826 is the determined cDNA sequence for the full-length
L551S clone 54298.

SEQ ID NO: 827 is the full-length amino acid sequence for L551S.

Tables 1-6 contain the sequence identifiers for SEQ ID NO:878-1664.

Table 1A

SEQ ID NO	CLONE IDENTIFIER	SEQ ID NO	CLONE IDENTIFIER
828	R0126:A02	869	R0126:D12
829	R0126:A03	870	R0126:E01
830	R0126:A05	871	R0126:E02
831	R0126:A06	872	R0126:E03
832	R0126:A08	873	R0126:E04
833	R0126:A09	874	R0126:E05
834	R0126:A10	875	R0126:E06
835	R0126:A11	876	R0126:E07
836	R0126:A12	877	R0126:E08
837	R0126:B01	878	R0126:E09
838	R0126:B03	879	R0126:E10
839	R0126:B04	880	R0126:E11
840	R0126:B05	881	R0126:E12
841	R0126:B06	882	R0126:F01
842	R0126:B07	883	R0126:F02
843	R0126:B08	884	R0126:F03
844	R0126:B09	885	R0126:F04
845	R0126:B11	886	R0126:F05
846	R0126:B12	887	R0126:F06
847	R0126:C01	888	R0126:F07
848	R0126:C02	889	R0126:F08
849	R0126:C03	890	R0126:F10
850	R0126:C05	891	R0126:F11
851	R0126:C06	892	R0126:F12
852	R0126:C07	893	R0126:G01
853	R0126:C08	894	R0126:G02
854	R0126:C09	895	R0126:G03
855	R0126:C10	896	R0126:G04
856	R0126:C11	897	R0126:G05
857	R0126:C12	898	R0126:G06
858	R0126:D01	899	R0126:G07
859	R0126:D02	900	R0126:G09
860	R0126:D03	901	R0126:G10
861	R0126:D04	902	R0126:G11
862	R0126:D05	903	R0126:G12
863	R0126:D06	904	R0126:H01
864	R0126:D07	905	R0126:H02
865	R0126:D08	906	R0126:H03
866	R0126:D09	907	R0126:H04

Table 1B

SEQ ID NO	CLONE IDENTIFIER	SEQ ID NO	CLONE IDENTIFIER
910	R0126:H07	951	R0127:D10
911	R0126:H09	952	R0127:D11
912	R0126:H10	953	R0127:D12
913	R0126:H11	954	R0127:E02
914	R0127:A02	955	R0127:E03
915	R0127:A05	956	R0127:E04
916	R0127:A06	957	R0127:E05
917	R0127:A07	958	R0127:E06
918	R0127:A08	959	R0127:E07
919	R0127:A09	960	R0127:E08
920	R0127:A10	961	R0127:E09
921	R0127:A11	962	R0127:E10
922	R0127:A12	963	R0127:E11
923	R0127:B01	964	R0127:F01
924	R0127:B03	965	R0127:F02
925	R0127:B04	966	R0127:F03
926	R0127:B05	967	R0127:F04
927	R0127:B06	968	R0127:F05
928	R0127:B07	969	R0127:F06
929	R0127:B08	970	R0127:F07
930	R0127:B09	971	R0127:F08
931	R0127:B10	972	R0127:F10
932	R0127:B11	973	R0127:F11
933	R0127:B12	974	R0127:F12
934	R0127:C01	975	R0127:G01
935	R0127:C03	976	R0127:G02
936	R0127:C04	977	R0127:G03
937	R0127:C05	978	R0127:G04
938	R0127:C07	979	R0127:G05
939	R0127:C08	980	R0127:G06
940	R0127:C09	981	R0127:G07
941	R0127:C10	982	R0127:G08
942	R0127:C11	983	R0127:G09
943	R0127:D01	984	R0127:G10
944	R0127:D02	985	R0127:G11
945	R0127:D03	986	R0127:G12
946	R0127:D04	987	R0127:H01
947	R0127:D05	988	R0127:H02
948	R0127:D06	989	R0127:H03

[illegible]

Table 1C

SEQ ID NO	CLONE IDENTIFIER	SEQ ID NO	CLONE IDENTIFIER
992	R0127:H06	1034	R0128:D11
993	R0127:H07	1035	R0128:D12
994	R0127:H08	1036	R0128:E01
995	R1027:H09	1037	R0128:E02
996	R1027:H10	1038	R0128:E03
997	R1027:H11	1039	R0128:E04
998	R1028:A02	1040	R0128:E05
999	R1028:A05	1041	R0128:E06
1000	R1028:A06	1042	R0128:E07
1001	R1028:A07	1043	R0128:E08
1002	R1028:A08	1044	R0128:E09
1003	R1028:A09	1045	R0128:E10
1004	R1028:A10	1046	R0128:E12
1005	R1028:B01	1047	R0128:F01
1006	R1028:B02	1048	R0128:F02
1007	R1028:B03	1049	R0128:F03
1008	R1028:B04	1050	R0128:F04
1009	R1028:B05	1051	R0128:F06
1010	R1028:B08	1052	R0128:F07
1011	R1028:B09	1053	R0128:F08
1012	R1028:B10	1054	R0128:F09
1013	R1028:B11	1055	R0128:F10
1014	R1028:B12	1056	R0128:F12
1015	R1028:C01	1057	R0128:G01
1016	R1028:C03	1058	R0128:G02
1017	R1028:C04	1059	R0128:G03
1018	R1028:C05	1060	R0128:G04
1019	R1028:C06	1061	R0128:G05
1020	R1028:C07	1062	R0128:G06
1021	R1028:C08	1063	R0128:G07
1022	R1028:C10	1064	R0128:G09
1023	R1028:C11	1065	R0128:G10
1024	R1028:C12	1066	R0128:G11
1025	R1028:D01	1067	R0128:G12
1026	R1028:D02	1068	R0128:H01
1027	R1028:D04	1069	R0128:H02
1028	R1028:D05	1070	R0128:H03
1029	R1028:D06	1071	R0128:H04
1030	R1028:D07	1072	R0128:H05
1031	R1028:D08	1073	R0128:H06

Table 1D

SEQ ID NO	CLONE IDENTIFIER	SEQ ID NO	CLONE IDENTIFIER
1076	R0128:H09	1118	R0130:D12
1077	R0128:H10	1119	R0130:E01
1078	R0128:H11	1120	R0130:E02
1079	R0130:A02	1121	R0130:E03
1080	R0130:A05	1122	R0130:E04
1081	R0130:A06	1123	R0130:E05
1082	R0130:A08	1124	R0130:E06
1083	R0130:A09	1125	R0130:E07
1084	R0130:A10	1126	R0130:E08
1085	R0130:A11	1127	R0130:E09
1086	R0130:A12	1128	R0130:E10
1087	R0130:B01	1129	R0130:E11
1088	R0130:B02	1130	R0130:E12
1089	R0130:B03	1131	R0130:F02
1090	R0130:B04	1132	R0130:F03
1091	R0130:B05	1133	R0130:F05
1092	R0130:B06	1134	R0130:F06
1093	R0130:B08	1135	R0130:F07
1094	R0130:B09	1136	R0130:F08
1095	R0130:B10	1137	R0130:F09
1096	R0130:B11	1138	R0130:F10
1097	R0130:B12	1139	R0130:F11
1098	R0130:C02	1140	R0130:F12
1099	R0130:C03	1141	R0130:G01
1100	R0130:C04	1142	R0130:G02
1101	R0130:C05	1143	R0130:G03
1102	R0130:C06	1144	R0130:G04
1103	R0130:C07	1145	R0130:G05
1104	R0130:C08	1146	R0130:G06
1105	R0130:C09	1147	R0130:G07
1106	R0130:C10	1148	R0130:G08
1107	R0130:C11	1149	R0130:G09
1108	R0130:C12	1150	R0130:G10
1109	R0130:D02	1151	R0130:G11
1110	R0130:D03	1152	R0130:G12
1111	R0130:D04	1153	R0130:H01
1112	R0130:D05	1154	R0130:H02
1113	R0130:D06	1155	R0130:H04
1114	R0130:D07	1156	R0130:H05
1115	R0130:D09	1157	R0130:H06

Table 1. Demographic characteristics of the study population	
Age (years)	50.0 ± 10.0
Gender (male/female)	100/100
Marital status (married/divorced/separated)	100/100/0
Education (years)	12.0 ± 2.0
Occupation (white/blue)	100/100
Income (USD/month)	1000.0 ± 200.0
Smoking status (smoker/nonsmoker)	50/50
Alcohol consumption (yes/no)	20/80
Family history of hypertension (yes/no)	30/70
Duration of hypertension (years)	5.0 ± 3.0
Current antihypertensive treatment (yes/no)	100/0
Medication (ACE inhibitor/CCB/β-blocker/diuretic)	100/0/0/0
Comorbidities (diabetes/cholesterol)	20/30
Quality of life (SF-36 score)	50.0 ± 10.0

Table 1E

SEQ ID NO	CLONE IDENTIFIER	SEQ ID NO	CLONE IDENTIFIER
1160	R0130:H09	1200	R0131:E01
1161	R0130:H10	1201	R0131:E02
1162	R0130:H11	1202	R0131:E03
1163	R0131:A02	1203	R0131:E04
1164	R0131:A05	1204	R0131:E06
1165	R0131:A06	1205	R0131:E07
1166	R0131:A07	1206	R0131:E08
1167	R0131:A08	1207	R0131:E10
1168	R0131:A09	1208	R0131:E11
1169	R0131:A11	1209	R0131:E12
1170	R0131:A12	1210	R0131:F02
1171	R0131:B02	1211	R0131:F04
1172	R0131:B03	1212	R0131:F05
1173	R0131:B04	1213	R0131:F06
1174	R0131:B05	1214	R0131:F07
1175	R0131:B07	1215	R0131:F08
1176	R0131:B08	1216	R0131:F09
1177	R0131:B09	1217	R0131:F10
1178	R0131:B10	1218	R0131:F11
1179	R0131:B11	1219	R0131:F12
1180	R0131:C01	1220	R0131:G01
1181	R0131:C02	1221	R0131:G02
1182	R0131:C03	1222	R0131:G03
1183	R0131:C04	1223	R0131:G04
1184	R0131:C06	1224	R0131:G05
1185	R0131:C07	1225	R0131:G06
1186	R0131:C08	1226	R0131:G07
1187	R0131:C10	1227	R0131:G08
1188	R0131:C11	1228	R0131:G09
1189	R0131:C12	1229	R0131:G10
1190	R0131:D02	1230	R0131:G11
1191	R0131:D03	1231	R0131:G12
1192	R0131:D04	1232	R0131:H01
1193	R0131:D05	1233	R0131:H02
1194	R0131:D06	1234	R0131:H05
1195	R0131:D07	1235	R0131:H06
1196	R0131:D09	1236	R0131:H07
1197	R0131:D10	1237	R0131:H08
1198	R0131:D11	1238	R0131:H09
1199	R0131:D12	1239	R0131:H11

Table 2:
Clone names for NSCLC-SQL1 and corresponding SEQ ID NOs

SEQ ID NO	CLONE IDENTIFIER	SEQ ID NO	CLONE IDENTIFIER
1240	Contig 54		
1241	Contig 55		
1242	Contig 57		
1243	Contig 58		
1244	Contig 60		
1245	Contig 62		
1246	Contig 63		
1247	Contig 64		
1248	Contig 65		
1249	Contig 66		
1250	Contig 67		
1251	Contig 68		
1252	Contig 69		
1253	Contig 70		
1254	Contig 71		
1255	Contig 72		
1256	Contig 73		
1257	Contig 74		
1258	Contig 75		
1259	Contig 77		
1260	Contig 78		
1261	Contig 79		
1262	Contig 80		
1263	Contig 81		
1264	Contig 83		
1265	Contig 84		
1266	Contig 86		
1267	Contig 87		
1268	Contig 88		
1269	Contig 89		
1270	Contig 90		
1271	Contig 91		
1272	Contig 92		
1273	Contig 94		
1274	Contig 95		
1275	Contig 96		
1276	Contig 97		
1277	Contig 98		

Table 3:
Clone names for NSCLC-SCLI and corresponding SEQ ID NOs

SEQ ID NO	CLONE IDENTIFIER	SEQ ID NO	CLONE IDENTIFIER
1280	Contig 38	1320	Contig 82
1281	Contig 39		
1282	Contig 40		
1283	Contig 41		
1284	Contig 42		
1285	Contig 43		
1286	Contig 44		
1287	Contig 45		
1288	Contig 46		
1289	Contig 47		
1290	Contig 48		
1291	Contig 49		
1292	Contig 51		
1293	Contig 52		
1294	Contig 53		
1295	Contig 54		
1296	Contig 55		
1297	Contig 56		
1298	Contig 57		
1299	Contig 58		
1300	Contig 59		
1301	Contig 60		
1302	Contig 62		
1303	Contig 63		
1304	Contig 64		
1305	Contig 65		
1306	Contig 66		
1307	Contig 67		
1308	Contig 68		
1309	Contig 69		
1310	Contig 70		
1311	Contig 72		
1312	Contig 73		
1313	Contig 75		
1314	Contig 76		
1315	Contig 77		
1316	Contig 78		
1317	Contig 79		

Table 4A:
Clone names for NSCLC-SCL3-SCL4 and corresponding SEQ ID NOs

SEQ ID NO	CLONE IDENTIFIER	SEQ ID NO	CLONE IDENTIFIER
1321	Contig 94	1363	Contig 136
1322	Contig 95	1364	Contig 137
1323	Contig 96	1365	Contig 138
1324	Contig 97	1366	Contig 139
1325	Contig 98	1367	Contig 140
1326	Contig 99	1368	Contig 141
1327	Contig 100	1369	Contig 142
1328	Contig 101	1370	Contig 143
1329	Contig 102	1371	Contig 144
1330	Contig 103	1372	Contig 145
1331	Contig 104	1373	Contig 146
1332	Contig 105	1374	Contig 147
1333	Contig 106	1375	Contig 148
1334	Contig 107	1376	Contig 149
1335	Contig 108	1377	Contig 150
1336	Contig 109	1378	Contig 151
1337	Contig 110	1379	Contig 152
1338	Contig 111	1380	Contig 153
1339	Contig 112	1381	Contig 154
1340	Contig 113	1382	Contig 155
1341	Contig 114	1383	Contig 156
1342	Contig 115	1384	Contig 157
1343	Contig 116	1385	Contig 158
1344	Contig 117	1386	Contig 159
1345	Contig 118	1387	Contig 160
1346	Contig 119	1388	Contig 161
1347	Contig 120	1389	Contig 162
1348	Contig 121	1390	Contig 163
1349	Contig 122	1391	Contig 164
1350	Contig 123	1392	Contig 165
1351	Contig 124	1393	Contig 166
1352	Contig 125	1394	Contig 167
1353	Contig 126	1395	Contig 168
1354	Contig 127	1396	Contig 169
1355	Contig 128	1397	Contig 170
1356	Contig 129	1398	Contig 171
1357	Contig 130	1399	Contig 172
1358	Contig 131	1400	Contig 173
1359	Contig 132	1401	Contig 174

SEQ ID NO	CLONE IDENTIFIER	SEQ ID NO	CLONE IDENTIFIER
1360	Contig 133	1402	Contig 175
1361	Contig 134	1403	Contig 176
1362	Contig 135		

Table 4B:
Clone names for NSCLC-SCL3-SCL4 and corresponding SEQ ID NOs

SEQ ID NO	CLONE IDENTIFIER	SEQ ID NO	CLONE IDENTIFIER
1404	Contig 177		
1405	Contig 178		
1406	Contig 179		
1407	Contig 180		
1408	Contig 181		
1409	Contig 182		
1410	Contig 183		
1411	Contig 184		
1412	Contig 185		
1413	Contig 186		
1414	Contig 187		

Table 5:
Clone names for SCLC-SQL1 and corresponding SEQ ID NOs

SEQ ID NO	CLONE IDENTIFIER	SEQ ID NO	CLONE IDENTIFIER
1415	Contig 17		
1416	Contig 18		
1417	Contig 20		
1418	Contig 23		
1419	Contig 24		
1420	Contig 25		
1421	Contig 26		
1422	Contig 27		
1423	Contig 28		
1424	Contig 29		
1425	Contig 30		
1426	Contig 31		
1427	Contig 20		
1428	Contig 21		
1429	Contig 22		
1430	Contig 23		
1431	Contig 24		
1432	Contig 25		
1433	Contig 26		
1434	Contig 27		
1435	Contig 28		
1436	Contig 29		
1437	Contig 30		
1438	Contig 31		
1439	Contig 32		
1440	Contig 33		
1441	Contig 34		
1442	Contig 35		
1443	Contig 36		
1444	Contig 37		
1445	Contig 38		

Table 6A:
Clone names for SCLC-SCL3-SCL4 and corresponding SEQ ID NOs

SEQ ID NO	CLONE IDENTIFIER	SEQ ID NO	CLONE IDENTIFIER
1446	Contig 116	1488	Contig 160
1447	Contig 117	1489	Contig 161
1448	Contig 118	1490	Contig 162
1449	Contig 119	1491	Contig 163
1450	Contig 120	1492	Contig 164
1451	Contig 122	1493	Contig 165
1452	Contig 123	1494	Contig 166
1453	Contig 124	1495	Contig 167
1454	Contig 125	1496	Contig 168
1455	Contig 126	1497	Contig 169
1456	Contig 127	1498	Contig 170
1457	Contig 128	1499	Contig 171
1458	Contig 129	1500	Contig 172
1459	Contig 130	1501	Contig 173
1460	Contig 131	1502	Contig 174
1461	Contig 132	1503	Contig 175
1462	Contig 133	1504	Contig 176
1463	Contig 135	1505	Contig 177
1464	Contig 136	1506	Contig 178
1465	Contig 137	1507	Contig 179
1466	Contig 138	1508	Contig 181
1467	Contig 139 (L985P)	1509	Contig 182
1468	Contig 140	1510	Contig 183
1469	Contig 141	1511	Contig 184
1470	Contig 142	1512	Contig 185
1471	Contig 143	1513	Contig 186
1472	Contig 144	1514	Contig 187
1473	Contig 145	1515	Contig 189
1474	Contig 146	1516	Contig 190
1475	Contig 147	1517	Contig 191
1476	Contig 148	1518	Contig 192
1477	Contig 149	1519	Contig 193
1478	Contig 150	1520	Contig 194
1479	Contig 151	1521	Contig 195
1480	Contig 152	1522	Contig 196
1481	Contig 153	1523	Contig 197
1482	Contig 154	1524	Contig 198
1483	Contig 155	1525	Contig 199
1484	Contig 156	1526	Contig 200

SEQ ID NO	CLONE IDENTIFIER	SEQ ID NO	CLONE IDENTIFIER
1485	Contig 157	1527	Contig 201
1486	Contig 158	1528	Contig 202
1487	Contig 159		

Contig 157

Table 6B:
Clone names for SCLC-SCL3-SCL4 and corresponding SEQ ID NOs

SEQ ID NO	CLONE IDENTIFIER	SEQ ID NO	CLONE IDENTIFIER
1529	Contig 203		
1530	Contig 204		
1531	Contig 205		
1532	Contig 206		
1533	Contig 207		
1534	Contig 208		
1535	Contig 209		
1536	Contig 210		
1537	Contig 211		
1538	Contig 212		
1539	Contig 213		
1540	Contig 214		
1541	Contig 215		
1542	Contig 216		
1543	Contig 217		
1544	Contig 218		
1545	Contig 219		
1546	Contig 220		
1547	Contig 221		
1548	Contig 222		
1549	Contig 223		
1550	Contig 224		
1551	Contig 225		
1552	Contig 226		
1553	Contig 227		
1554	Contig 228		
1555	Contig 229		
1556	Contig 230		
1557	Contig 231		
1558	Contig 232		
1559	Contig 233		
1560	Contig 234		
1561	Contig 235		
1562	Contig 236		
1563	Contig 237		

Table 7.

SEQ ID NO	CLONE IDENTIFIER	SEQ ID NO	CLONE IDENTIFIER
1564	R0124:E05	1609	R0129:D09
1565	R0124:E06	1610	R0129:D10
1566	R0124:E08	1611	R0129:D11
1567	R0124:F07	1612	R0129:E02
1568	R0124:F08	1613	R0129:E03
1569	R0124:F09	1614	R0129:E04
1570	R0124:G04	1615	R0129:E05
1571	R0129:A02	1616	R0129:E06
1572	R0129:A03	1617	R0129:E07
1573	R0129:A06	1618	R0129:E08
1574	R0129:A07	1619	R0129:E09
1575	R0129:A08	1620	R0129:E11
1576	R0129:A09	1621	R0129:E12
1577	R0129:A10	1622	R0129:F01
1578	R0129:A11	1623	R0129:F02
1579	R0129:A12	1624	R0129:F03
1580	R0129:B02	1625	R0129:F04
1581	R0129:B03	1626	R0129:F06
1582	R0129:B04	1627	R0129:F07
1583	R0129:B05	1628	R0129:F08
1584	R0129:B06	1629	R0129:F09
1585	R0129:B07	1630	R0129:F10
1586	R0129:B08	1631	R0129:F11
1587	R0129:B09	1632	R0129:F12
1588	R0129:B10	1633	R0129:G01
1589	R0129:B11	1634	R0129:G02
1590	R0129:B12	1635	R0129:G03
1591	R0129:C01	1636	R0129:G04
1592	R0129:C02	1637	R0129:G05
1593	R0129:C03	1638	R0129:G06
1594	R0129:C04	1639	R0129:G07
1595	R0129:C06	1640	R0129:G08
1596	R0129:C07	1641	R0129:G09
1597	R0129:C08	1642	R0129:G10
1598	R0129:C09	1643	R0129:G11
1599	R0129:C10	1644	R0129:G12
1600	R0129:C11	1645	R0129:H01
1601	R0129:C12	1646	R0129:H02
1602	R0129:D01	1647	R0129:H03
1603	R0129:D03	1648	R0129:H04

SEQ ID NO	CLONE IDENTIFIER	SEQ ID NO	CLONE IDENTIFIER
1604	R0129:D04	1649	R0129:H05
1605	R0129:D05	1650	R0129:H08
1606	R0129:D06	1651	R0129:H09
1607	R0129:D07	1652	R0129:H10
1608	R0129:D08	1653	R0129:H11

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Table 8.

SEQ ID NO	CLONE IDENTIFIER	SEQ ID NO	CLONE IDENTIFIER
1654	26484		
1655	26496		
1656	26517		
1657	26531		
1658	26022		
1659	26026		
1660	26810		
1661	26815		
1662	26869		
1663	26883		
1664	26902		

SEQ ID NO:1667 is the protein sequence of expressed recombinant

5 L7548S.

SEQ ID NO:1668 is the cDNA sequence of expressed recombinant L7548S.

SEQ ID NO:1669 is the extended cDNA sequence of clone #18971

(L801P).

10 SEQ ID NO:1670 is the amino acid sequence of open reading frame ORF4
encoded by SEQ ID NO:1669.

SEQ ID NO:1671 is the amino acid sequence of open reading frame ORF5
encoded by SEQ ID NO:1669.

SEQ ID NO:1672 is the amino acid sequence of open reading frame ORF6
encoded by SEQ ID NO:1669.

15 SEQ ID NO:1673 is the amino acid sequence of open reading frame ORF7
encoded by SEQ ID NO:1669.

SEQ ID NO:1674 is the amino acid sequence of open reading frame ORF8
encoded by SEQ ID NO:1669.

20 SEQ ID NO:1675 is the amino acid sequence of open reading frame ORF9
encoded by SEQ ID NO:1669.

SEQ ID NO:1676 is the extended cDNA for contig 139 (SEQ ID NO:1467),
also known as L985P.

SEQ ID NO:1677 is the L985P amino acid sequence encoded by SEQ ID NO: 1676.

SEQ ID NO: 1678 is the amino acid sequence of open reading frame ORF5X of SEQ ID NO:1669.

- 5 SEQ ID NO: 1679 is the amino acid sequence of an open reading frame for contig 139 (SEQ ID NO:1467).

DETAILED DESCRIPTION OF THE INVENTION

As noted above, the present invention is generally directed to compositions and methods for using the compositions, for example in the therapy and diagnosis of cancer, such as lung cancer. Certain illustrative compositions described herein include lung tumor polypeptides, polynucleotides encoding such polypeptides, binding agents such as antibodies, antigen presenting cells (APCs) and/or immune system cells (*e.g.*, T cells). A "lung tumor protein," as the term is used herein, refers generally to a protein that is expressed in lung tumor cells at a level that is at least two fold, and preferably at least five fold, greater than the level of expression in a normal tissue, as determined using a representative assay provided herein. Certain lung tumor proteins are tumor proteins that react detectably (within an immunoassay, such as an ELISA or Western blot) with antisera of a patient afflicted with lung cancer.

20 Therefore, in accordance with the above, and as described further below, the present invention provides illustrative polynucleotide compositions having sequences set forth in SEQ ID NO: 1, 11-13, 15, 20, 23-27, 29, 30, 33, 34, 39, 41, 43-46, 51, 52, 57, 58, 60, 62, 65-67, 69-71, 74, 76, 79, 80, 84, 86, 89-92, 95, 97, 98, 101, 110, 111, 113-119, 121-128, 130-134, 136, 138, 139, 141, 143, 146-151, 153, 154, 157-160, 162-164, 167-178, 180, 181, 183, 186-190, 192, 193, 195-220, 224, 226-231, 234, 236, 237, 240, 241, 244-246, 248, 254, 255, 261, 262, 266, 270, 275, 280, 282, 283, 288, 289, 290, 292, 295, 301, 303, 304, 309, 311, 341-782, 784, 785, 790, 792, 794, 796, 800-804, 807, 808, 810-826, 1240, 1243, 1247, 1269, 1272, 1280, 1283, 1285, 1286, 1289, 1300, 1309, 1318, 1319,

1327, 1335, 1339, 1346, 1359, 1369, 1370, 1371, 1393, 1398, 1405, 1408, 1413, 1414, 1417, 1422, 1429, 1432, 1435, 1436, 1438-1442, 1447, 1450, 1453, 1463, 1467, 1470, 1473, 1475, 1482, 1486, 1491-1494, 1501, 1505, 1506, 1514-1517, 1520, 1522, 1524, 1535, 1538, 1542, 1543, 1547, 1554, 1557, 1559, 1561, 1563 and 1669 illustrative
 5 polypeptide compositions having amino acid sequences set forth in SEQ ID NO: 786, 787, 791, 793, 795, 797-799, 806, 809, 827, 1670-1675 and 1677-1679, antibody compositions capable of binding such polypeptides, and numerous additional embodiments employing such compositions, for example in the detection, diagnosis and/or therapy of human lung cancer.

10 POLYNUCLEOTIDE COMPOSITIONS

As used herein, the terms "DNA segment" and "polynucleotide" refer to a DNA molecule that has been isolated free of total genomic DNA of a particular species. Therefore, a DNA segment encoding a polypeptide refers to a DNA segment that contains one or more coding sequences yet is substantially isolated away from, or purified free from,
 15 total genomic DNA of the species from which the DNA segment is obtained. Included within the terms "DNA segment" and "polynucleotide" are DNA segments and smaller fragments of such segments, and also recombinant vectors, including, for example, plasmids, cosmids, phagemids, phage, viruses, and the like.

As will be understood by those skilled in the art, the DNA segments of this
 20 invention can include genomic sequences, extra-genomic and plasmid-encoded sequences and smaller engineered gene segments that express, or may be adapted to express, proteins, polypeptides, peptides and the like. Such segments may be naturally isolated, or modified synthetically by the hand of man.

"Isolated," as used herein, means that a polynucleotide is substantially away
 25 from other coding sequences, and that the DNA segment does not contain large portions of unrelated coding DNA, such as large chromosomal fragments or other functional genes or polypeptide coding regions. Of course, this refers to the DNA segment as originally

isolated, and does not exclude genes or coding regions later added to the segment by the hand of man.

As will be recognized by the skilled artisan, polynucleotides may be single-stranded (coding or antisense) or double-stranded, and may be DNA (genomic, cDNA or synthetic) or RNA molecules. RNA molecules include HnRNA molecules, which contain introns and correspond to a DNA molecule in a one-to-one manner, and mRNA molecules, which do not contain introns. Additional coding or non-coding sequences may, but need not, be present within a polynucleotide of the present invention, and a polynucleotide may, but need not, be linked to other molecules and/or support materials.

Polynucleotides may comprise a native sequence (*i.e.*, an endogenous sequence that encodes a lung tumor protein or a portion thereof) or may comprise a variant, or a biological or antigenic functional equivalent of such a sequence. Polynucleotide variants may contain one or more substitutions, additions, deletions and/or insertions, as further described below, preferably such that the immunogenicity of the encoded polypeptide is not diminished, relative to a native tumor protein. The effect on the immunogenicity of the encoded polypeptide may generally be assessed as described herein. The term “variants” also encompasses homologous genes of xenogenic origin.

When comparing polynucleotide or polypeptide sequences, two sequences are said to be “identical” if the sequence of nucleotides or amino acids in the two sequences is the same when aligned for maximum correspondence, as described below. Comparisons between two sequences are typically performed by comparing the sequences over a comparison window to identify and compare local regions of sequence similarity. A “comparison window” as used herein, refers to a segment of at least about 20 contiguous positions, usually 30 to about 75, 40 to about 50, in which a sequence may be compared to a reference sequence of the same number of contiguous positions after the two sequences are optimally aligned.

Optimal alignment of sequences for comparison may be conducted using the Megalign program in the Lasergene suite of bioinformatics software (DNASTAR, Inc., Madison, WI), using default parameters. This program embodies several alignment

- schemes described in the following references: Dayhoff, M.O. (1978) A model of evolutionary change in proteins – Matrices for detecting distant relationships. In Dayhoff, M.O. (ed.) *Atlas of Protein Sequence and Structure*, National Biomedical Research Foundation, Washington DC Vol. 5, Suppl. 3, pp. 345-358; Hein J. (1990) Unified
- 5 Approach to Alignment and Phylogenesis pp. 626-645 *Methods in Enzymology* vol. 183, Academic Press, Inc., San Diego, CA; Higgins, D.G. and Sharp, P.M. (1989) *CABIOS* 5:151-153; Myers, E.W. and Muller W. (1988) *CABIOS* 4:11-17; Robinson, E.D. (1971) *Comb. Theor* 11:105; Santou, N. Nes, M. (1987) *Mol. Biol. Evol.* 4:406-425; Sneath, P.H.A. and Sokal, R.R. (1973) *Numerical Taxonomy – the Principles and Practice of*
- 10 *Numerical Taxonomy*, Freeman Press, San Francisco, CA; Wilbur, W.J. and Lipman, D.J. (1983) *Proc. Natl. Acad., Sci. USA* 80:726-730.

Alternatively, optimal alignment of sequences for comparison may be conducted by the local identity algorithm of Smith and Waterman (1981) *Add. APL. Math* 2:482, by the identity alignment algorithm of Needleman and Wunsch (1970) *J. Mol. Biol.*

15 48:443, by the search for similarity methods of Pearson and Lipman (1988) *Proc. Natl. Acad. Sci. USA* 85: 2444, by computerized implementations of these algorithms (GAP, BESTFIT, BLAST, FASTA, and TFASTA in the Wisconsin Genetics Software Package, Genetics Computer Group (GCG), 575 Science Dr., Madison, WI), or by inspection.

One preferred example of algorithms that are suitable for determining

20 percent sequence identity and sequence similarity are the BLAST and BLAST 2.0 algorithms, which are described in Altschul *et al.* (1977) *Nucl. Acids Res.* 25:3389-3402 and Altschul *et al.* (1990) *J. Mol. Biol.* 215:403-410, respectively. BLAST and BLAST 2.0 can be used, for example with the parameters described herein, to determine percent sequence identity for the polynucleotides and polypeptides of the invention. Software for

25 performing BLAST analyses is publicly available through the National Center for Biotechnology Information. In one illustrative example, cumulative scores can be calculated using, for nucleotide sequences, the parameters M (reward score for a pair of matching residues; always >0) and N (penalty score for mismatching residues; always <0). For amino acid sequences, a scoring matrix can be used to calculate the cumulative score.

Extension of the word hits in each direction are halted when: the cumulative alignment score falls off by the quantity X from its maximum achieved value; the cumulative score goes to zero or below, due to the accumulation of one or more negative-scoring residue alignments; or the end of either sequence is reached. The BLAST algorithm parameters W, T and X determine the sensitivity and speed of the alignment. The BLASTN program (for nucleotide sequences) uses as defaults a wordlength (W) of 11, and expectation (E) of 10, and the BLOSUM62 scoring matrix (see Henikoff and Henikoff (1989) *Proc. Natl. Acad. Sci. USA* 89:10915) alignments, (B) of 50, expectation (E) of 10, M=5, N=-4 and a comparison of both strands.

Preferably, the "percentage of sequence identity" is determined by comparing two optimally aligned sequences over a window of comparison of at least 20 positions, wherein the portion of the polynucleotide or polypeptide sequence in the comparison window may comprise additions or deletions (*i.e.*, gaps) of 20 percent or less, usually 5 to 15 percent, or 10 to 12 percent, as compared to the reference sequences (which does not comprise additions or deletions) for optimal alignment of the two sequences. The percentage is calculated by determining the number of positions at which the identical nucleic acid bases or amino acid residue occurs in both sequences to yield the number of matched positions, dividing the number of matched positions by the total number of positions in the reference sequence (*i.e.*, the window size) and multiplying the results by 100 to yield the percentage of sequence identity.

Therefore, the present invention encompasses polynucleotide and polypeptide sequences having substantial identity to the sequences disclosed herein, for example those comprising at least 50% sequence identity, preferably at least 55%, 60%, 65%, 70%, 75%, 80%, 85%, 90%, 95%, 96%, 97%, 98%, or 99% or higher, sequence identity compared to a polynucleotide or polypeptide sequence of this invention using the methods described herein, (*e.g.*, BLAST analysis using standard parameters, as described below). One skilled in this art will recognize that these values can be appropriately adjusted to determine corresponding identity of proteins encoded by two nucleotide

sequences by taking into account codon degeneracy, amino acid similarity, reading frame positioning and the like.

In additional embodiments, the present invention provides isolated polynucleotides and polypeptides comprising various lengths of contiguous stretches of
 5 sequence identical to or complementary to one or more of the sequences disclosed herein. For example, polynucleotides are provided by this invention that comprise at least about 15, 20, 30, 40, 50, 75, 100, 150, 200, 300, 400, 500 or 1000 or more contiguous nucleotides of one or more of the sequences disclosed herein as well as all intermediate lengths there between. It will be readily understood that "intermediate lengths", in this context, means
 10 any length between the quoted values, such as 16, 17, 18, 19, *etc.*; 21, 22, 23, *etc.*; 30, 31, 32, *etc.*; 50, 51, 52, 53, *etc.*; 100, 101, 102, 103, *etc.*; 150, 151, 152, 153, *etc.*; including all integers through 200-500; 500-1,000, and the like.

The polynucleotides of the present invention, or fragments thereof, regardless of the length of the coding sequence itself, may be combined with other DNA
 15 sequences, such as promoters, polyadenylation signals, additional restriction enzyme sites, multiple cloning sites, other coding segments, and the like, such that their overall length may vary considerably. It is therefore contemplated that a nucleic acid fragment of almost any length may be employed, with the total length preferably being limited by the ease of preparation and use in the intended recombinant DNA protocol. For example, illustrative
 20 DNA segments with total lengths of about 10,000, about 5000, about 3000, about 2,000, about 1,000, about 500, about 200, about 100, about 50 base pairs in length, and the like, (including all intermediate lengths) are contemplated to be useful in many implementations of this invention.

In other embodiments, the present invention is directed to polynucleotides
 25 that are capable of hybridizing under moderately stringent conditions to a polynucleotide sequence provided herein, or a fragment thereof, or a complementary sequence thereof. Hybridization techniques are well known in the art of molecular biology. For purposes of illustration, suitable moderately stringent conditions for testing the hybridization of a polynucleotide of this invention with other polynucleotides include prewashing in a

solution of 5 X SSC, 0.5% SDS, 1.0 mM EDTA (pH 8.0); hybridizing at 50°C-65°C, 5 X SSC, overnight; followed by washing twice at 65°C for 20 minutes with each of 2X, 0.5X and 0.2X SSC containing 0.1% SDS.

Moreover, it will be appreciated by those of ordinary skill in the art that, as a result of the degeneracy of the genetic code, there are many nucleotide sequences that encode a polypeptide as described herein. Some of these polynucleotides bear minimal homology to the nucleotide sequence of any native gene. Nonetheless, polynucleotides that vary due to differences in codon usage are specifically contemplated by the present invention. Further, alleles of the genes comprising the polynucleotide sequences provided herein are within the scope of the present invention. Alleles are endogenous genes that are altered as a result of one or more mutations, such as deletions, additions and/or substitutions of nucleotides. The resulting mRNA and protein may, but need not, have an altered structure or function. Alleles may be identified using standard techniques (such as hybridization, amplification and/or database sequence comparison).

15 **PROBES AND PRIMERS**

In other embodiments of the present invention, the polynucleotide sequences provided herein can be advantageously used as probes or primers for nucleic acid hybridization. As such, it is contemplated that nucleic acid segments that comprise a sequence region of at least about 15 nucleotide long contiguous sequence that has the same sequence as, or is complementary to, a 15 nucleotide long contiguous sequence disclosed herein will find particular utility. Longer contiguous identical or complementary sequences, *e.g.*, those of about 20, 30, 40, 50, 100, 200, 500, 1000 (including all intermediate lengths) and even up to full length sequences will also be of use in certain embodiments.

25 The ability of such nucleic acid probes to specifically hybridize to a sequence of interest will enable them to be of use in detecting the presence of complementary sequences in a given sample. However, other uses are also envisioned,

such as the use of the sequence information for the preparation of mutant species primers, or primers for use in preparing other genetic constructions.

Polynucleotide molecules having sequence regions consisting of contiguous nucleotide stretches of 10-14, 15-20, 30, 50, or even of 100-200 nucleotides or so (including intermediate lengths as well), identical or complementary to a polynucleotide sequence disclosed herein, are particularly contemplated as hybridization probes for use in, *e.g.*, Southern and Northern blotting. This would allow a gene product, or fragment thereof, to be analyzed, both in diverse cell types and also in various bacterial cells. The total size of fragment, as well as the size of the complementary stretch(es), will ultimately depend on the intended use or application of the particular nucleic acid segment. Smaller fragments will generally find use in hybridization embodiments, wherein the length of the contiguous complementary region may be varied, such as between about 15 and about 100 nucleotides, but larger contiguous complementarity stretches may be used, according to the length complementary sequences one wishes to detect.

The use of a hybridization probe of about 15-25 nucleotides in length allows the formation of a duplex molecule that is both stable and selective. Molecules having contiguous complementary sequences over stretches greater than 15 bases in length are generally preferred, though, in order to increase stability and selectivity of the hybrid, and thereby improve the quality and degree of specific hybrid molecules obtained. One will generally prefer to design nucleic acid molecules having gene-complementary stretches of 15 to 25 contiguous nucleotides, or even longer where desired.

Hybridization probes may be selected from any portion of any of the sequences disclosed herein. All that is required is to review the sequence set forth in SEQ ID NO: 1, 11-13, 15, 20, 23-27, 29, 30, 33, 34, 39, 41, 43-46, 51, 52, 57, 58, 60, 62, 65-67, 69-71, 74, 76, 79, 80, 84, 86, 89-92, 95, 97, 98, 101, 110, 111, 113-119, 121-128, 130-134, 136, 138, 139, 141, 143, 146-151, 153, 154, 157-160, 162-164, 167-178, 180, 181, 183, 186-190, 192, 193, 195-220, 224, 226-231, 234, 236, 237, 240, 241, 244-246, 248, 254, 255, 261, 262, 266, 270, 275, 280, 282, 283, 288, 289, 290, 292, 295, 301, 303, 304, 309, 311, 341-782, 784, 785, 790, 792, 794, 796, 800-804, 807, 808, 810-826, 828-1664, 1669

and 1676, or to any continuous portion of the sequence, from about 15-25 nucleotides in length up to and including the full length sequence, that one wishes to utilize as a probe or primer. The choice of probe and primer sequences may be governed by various factors. For example, one may wish to employ primers from towards the termini of the total
5 sequence.

Small polynucleotide segments or fragments may be readily prepared by, for example, directly synthesizing the fragment by chemical means, as is commonly practiced using an automated oligonucleotide synthesizer. Also, fragments may be obtained by application of nucleic acid reproduction technology, such as the PCR™ technology of U. S.
10 Patent 4,683,202 (incorporated herein by reference), by introducing selected sequences into recombinant vectors for recombinant production, and by other recombinant DNA techniques generally known to those of skill in the art of molecular biology.

The nucleotide sequences of the invention may be used for their ability to selectively form duplex molecules with complementary stretches of the entire gene or gene
15 fragments of interest. Depending on the application envisioned, one will typically desire to employ varying conditions of hybridization to achieve varying degrees of selectivity of probe towards target sequence. For applications requiring high selectivity, one will typically desire to employ relatively stringent conditions to form the hybrids, *e.g.*, one will select relatively low salt and/or high temperature conditions, such as provided by a salt
20 concentration of from about 0.02 M to about 0.15 M salt at temperatures of from about 50°C to about 70°C. Such selective conditions tolerate little, if any, mismatch between the probe and the template or target strand, and would be particularly suitable for isolating related sequences.

Of course, for some applications, for example, where one desires to prepare
25 mutants employing a mutant primer strand hybridized to an underlying template, less stringent (reduced stringency) hybridization conditions will typically be needed in order to allow formation of the heteroduplex. In these circumstances, one may desire to employ salt conditions such as those of from about 0.15 M to about 0.9 M salt, at temperatures ranging from about 20°C to about 55°C. Cross-hybridizing species can thereby be readily

identified as positively hybridizing signals with respect to control hybridizations. In any case, it is generally appreciated that conditions can be rendered more stringent by the addition of increasing amounts of formamide, which serves to destabilize the hybrid duplex in the same manner as increased temperature. Thus, hybridization conditions can be readily manipulated, and thus will generally be a method of choice depending on the desired results.

POLYNUCLEOTIDE IDENTIFICATION AND CHARACTERIZATION

Polynucleotides may be identified, prepared and/or manipulated using any of a variety of well established techniques. For example, a polynucleotide may be identified, as described in more detail below, by screening a microarray of cDNAs for tumor-associated expression (*i.e.*, expression that is at least two fold greater in a tumor than in normal tissue, as determined using a representative assay provided herein). Such screens may be performed, for example, using a Synteni microarray (Palo Alto, CA) according to the manufacturer's instructions (and essentially as described by Schena *et al.*, *Proc. Natl. Acad. Sci. USA* 93:10614-10619, 1996 and Heller *et al.*, *Proc. Natl. Acad. Sci. USA* 94:2150-2155, 1997). Alternatively, polynucleotides may be amplified from cDNA prepared from cells expressing the proteins described herein, such as lung tumor cells. Such polynucleotides may be amplified via polymerase chain reaction (PCR). For this approach, sequence-specific primers may be designed based on the sequences provided herein, and may be purchased or synthesized.

An amplified portion of a polynucleotide of the present invention may be used to isolate a full length gene from a suitable library (*e.g.*, a lung tumor cDNA library) using well known techniques. Within such techniques, a library (cDNA or genomic) is screened using one or more polynucleotide probes or primers suitable for amplification. Preferably, a library is size-selected to include larger molecules. Random primed libraries may also be preferred for identifying 5' and upstream regions of genes. Genomic libraries are preferred for obtaining introns and extending 5' sequences.

For hybridization techniques, a partial sequence may be labeled (*e.g.*, by nick-translation or end-labeling with ^{32}P) using well known techniques. A bacterial or bacteriophage library is then generally screened by hybridizing filters containing denatured bacterial colonies (or lawns containing phage plaques) with the labeled probe (*see*

5 Sambrook *et al.*, *Molecular Cloning: A Laboratory Manual*, Cold Spring Harbor Laboratories, Cold Spring Harbor, NY, 1989). Hybridizing colonies or plaques are selected and expanded, and the DNA is isolated for further analysis. cDNA clones may be analyzed to determine the amount of additional sequence by, for example, PCR using a

10 partial sequences may be generated to identify one or more overlapping clones. The complete sequence may then be determined using standard techniques, which may involve generating a series of deletion clones. The resulting overlapping sequences can then assembled into a single contiguous sequence. A full length cDNA molecule can be generated by ligating suitable fragments, using well known techniques.

15 Alternatively, there are numerous amplification techniques for obtaining a full length coding sequence from a partial cDNA sequence. Within such techniques, amplification is generally performed via PCR. Any of a variety of commercially available kits may be used to perform the amplification step. Primers may be designed using, for example, software well known in the art. Primers are preferably 22-30 nucleotides in

20 length, have a GC content of at least 50% and anneal to the target sequence at temperatures of about 68°C to 72°C. The amplified region may be sequenced as described above, and overlapping sequences assembled into a contiguous sequence.

One such amplification technique is inverse PCR (*see* Triglia *et al.*, *Nucl. Acids Res.* 16:8186, 1988), which uses restriction enzymes to generate a fragment in the

25 known region of the gene. The fragment is then circularized by intramolecular ligation and used as a template for PCR with divergent primers derived from the known region. Within an alternative approach, sequences adjacent to a partial sequence may be retrieved by amplification with a primer to a linker sequence and a primer specific to a known region. The amplified sequences are typically subjected to a second round of amplification with the

same linker primer and a second primer specific to the known region. A variation on this procedure, which employs two primers that initiate extension in opposite directions from the known sequence, is described in WO 96/38591. Another such technique is known as "rapid amplification of cDNA ends" or RACE. This technique involves the use of an
 5 internal primer and an external primer, which hybridizes to a polyA region or vector sequence, to identify sequences that are 5' and 3' of a known sequence. Additional techniques include capture PCR (Lagerstrom *et al.*, *PCR Methods Applic.* 1:111-19, 1991) and walking PCR (Parker *et al.*, *Nucl. Acids. Res.* 19:3055-60, 1991). Other methods employing amplification may also be employed to obtain a full length cDNA sequence.

10 In certain instances, it is possible to obtain a full length cDNA sequence by analysis of sequences provided in an expressed sequence tag (EST) database, such as that available from GenBank. Searches for overlapping ESTs may generally be performed using well known programs (*e.g.*, NCBI BLAST searches), and such ESTs may be used to generate a contiguous full length sequence. Full length DNA sequences may also be
 15 obtained by analysis of genomic fragments.

POLYNUCLEOTIDE EXPRESSION IN HOST CELLS

In other embodiments of the invention, polynucleotide sequences or fragments thereof which encode polypeptides of the invention, or fusion proteins or functional equivalents thereof, may be used in recombinant DNA molecules to direct
 20 expression of a polypeptide in appropriate host cells. Due to the inherent degeneracy of the genetic code, other DNA sequences that encode substantially the same or a functionally equivalent amino acid sequence may be produced and these sequences may be used to clone and express a given polypeptide.

As will be understood by those of skill in the art, it may be advantageous in
 25 some instances to produce polypeptide-encoding nucleotide sequences possessing non-naturally occurring codons. For example, codons preferred by a particular prokaryotic or eukaryotic host can be selected to increase the rate of protein expression or to produce a

recombinant RNA transcript having desirable properties, such as a half-life which is longer than that of a transcript generated from the naturally occurring sequence.

Moreover, the polynucleotide sequences of the present invention can be engineered using methods generally known in the art in order to alter polypeptide encoding sequences for a variety of reasons, including but not limited to, alterations which modify the cloning, processing, and/or expression of the gene product. For example, DNA shuffling by random fragmentation and PCR reassembly of gene fragments and synthetic oligonucleotides may be used to engineer the nucleotide sequences. In addition, site-directed mutagenesis may be used to insert new restriction sites, alter glycosylation patterns, change codon preference, produce splice variants, or introduce mutations, and so forth.

In another embodiment of the invention, natural, modified, or recombinant nucleic acid sequences may be ligated to a heterologous sequence to encode a fusion protein. For example, to screen peptide libraries for inhibitors of polypeptide activity, it may be useful to encode a chimeric protein that can be recognized by a commercially available antibody. A fusion protein may also be engineered to contain a cleavage site located between the polypeptide-encoding sequence and the heterologous protein sequence, so that the polypeptide may be cleaved and purified away from the heterologous moiety.

Sequences encoding a desired polypeptide may be synthesized, in whole or in part, using chemical methods well known in the art (see Caruthers, M. H. *et al.* (1980) *Nucl. Acids Res. Symp. Ser.* 215-223, Horn, T. *et al.* (1980) *Nucl. Acids Res. Symp. Ser.* 225-232). Alternatively, the protein itself may be produced using chemical methods to synthesize the amino acid sequence of a polypeptide, or a portion thereof. For example, peptide synthesis can be performed using various solid-phase techniques (Roberge, J. Y. *et al.* (1995) *Science* 269:202-204) and automated synthesis may be achieved, for example, using the ABI 431A Peptide Synthesizer (Perkin Elmer, Palo Alto, CA).

A newly synthesized peptide may be substantially purified by preparative high performance liquid chromatography (*e.g.*, Creighton, T. (1983) *Proteins, Structures and Molecular Principles*, WH Freeman and Co., New York, N.Y.) or other comparable

techniques available in the art. The composition of the synthetic peptides may be confirmed by amino acid analysis or sequencing (*e.g.*, the Edman degradation procedure). Additionally, the amino acid sequence of a polypeptide, or any part thereof, may be altered during direct synthesis and/or combined using chemical methods with sequences from
 5 other proteins, or any part thereof, to produce a variant polypeptide.

In order to express a desired polypeptide, the nucleotide sequences encoding the polypeptide, or functional equivalents, may be inserted into appropriate expression vector, *i.e.*, a vector which contains the necessary elements for the transcription and translation of the inserted coding sequence. Methods which are well known to those
 10 skilled in the art may be used to construct expression vectors containing sequences encoding a polypeptide of interest and appropriate transcriptional and translational control elements. These methods include *in vitro* recombinant DNA techniques, synthetic techniques, and *in vivo* genetic recombination. Such techniques are described in Sambrook, J. *et al.* (1989) *Molecular Cloning, A Laboratory Manual*, Cold Spring Harbor Press, Plainview, N.Y., and Ausubel, F. M. *et al.* (1989) *Current Protocols in Molecular Biology*,
 15 John Wiley & Sons, New York, N.Y.

A variety of expression vector/host systems may be utilized to contain and express polynucleotide sequences. These include, but are not limited to, microorganisms such as bacteria transformed with recombinant bacteriophage, plasmid, or cosmid DNA
 20 expression vectors; yeast transformed with yeast expression vectors; insect cell systems infected with virus expression vectors (*e.g.*, baculovirus); plant cell systems transformed with virus expression vectors (*e.g.*, cauliflower mosaic virus, CaMV; tobacco mosaic virus, TMV) or with bacterial expression vectors (*e.g.*, Ti or pBR322 plasmids); or animal cell systems.

25 The "control elements" or "regulatory sequences" present in an expression vector are those non-translated regions of the vector--enhancers, promoters, 5' and 3' untranslated regions--which interact with host cellular proteins to carry out transcription and translation. Such elements may vary in their strength and specificity. Depending on the vector system and host utilized, any number of suitable transcription and translation

elements, including constitutive and inducible promoters, may be used. For example, when cloning in bacterial systems, inducible promoters such as the hybrid lacZ promoter of the PBLUESCRIPT phagemid (Stratagene, La Jolla, Calif.) or PSFORT1 plasmid (Gibco BRL, Gaithersburg, MD) and the like may be used. In mammalian cell systems, promoters from

5 mammalian genes or from mammalian viruses are generally preferred. If it is necessary to generate a cell line that contains multiple copies of the sequence encoding a polypeptide, vectors based on SV40 or EBV may be advantageously used with an appropriate selectable marker.

In bacterial systems, a number of expression vectors may be selected

10 depending upon the use intended for the expressed polypeptide. For example, when large quantities are needed, for example for the induction of antibodies, vectors which direct high level expression of fusion proteins that are readily purified may be used. Such vectors include, but are not limited to, the multifunctional *E. coli* cloning and expression vectors such as BLUESCRIPT (Stratagene), in which the sequence encoding the polypeptide of

15 interest may be ligated into the vector in frame with sequences for the amino-terminal Met and the subsequent 7 residues of β -galactosidase so that a hybrid protein is produced; pIN vectors (Van Heeke, G. and S. M. Schuster (1989) *J. Biol. Chem.* 264:5503-5509); and the like. pGEX Vectors (Promega, Madison, Wis.) may also be used to express foreign polypeptides as fusion proteins with glutathione S-transferase (GST). In general, such

20 fusion proteins are soluble and can easily be purified from lysed cells by adsorption to glutathione-agarose beads followed by elution in the presence of free glutathione. Proteins made in such systems may be designed to include heparin, thrombin, or factor XA protease cleavage sites so that the cloned polypeptide of interest can be released from the GST moiety at will.

25 In the yeast, *Saccharomyces cerevisiae*, a number of vectors containing constitutive or inducible promoters such as alpha factor, alcohol oxidase, and PGH may be used. For reviews, see Ausubel *et al.* (supra) and Grant *et al.* (1987) *Methods Enzymol.* 153:516-544.

In cases where plant expression vectors are used, the expression of sequences encoding polypeptides may be driven by any of a number of promoters. For example, viral promoters such as the 35S and 19S promoters of CaMV may be used alone or in combination with the omega leader sequence from TMV (Takamatsu, N. (1987) *EMBO J.* 6:307-311. Alternatively, plant promoters such as the small subunit of RUBISCO or heat shock promoters may be used (Coruzzi, G. *et al.* (1984) *EMBO J.* 3:1671-1680; Broglie, R. *et al.* (1984) *Science* 224:838-843; and Winter, J. *et al.* (1991) *Results Probl. Cell Differ.* 17:85-105). These constructs can be introduced into plant cells by direct DNA transformation or pathogen-mediated transfection. Such techniques are described in a number of generally available reviews (see, for example, Hobbs, S. or Murry, L. E. in McGraw Hill Yearbook of Science and Technology (1992) McGraw Hill, New York, N.Y.; pp. 191-196).

An insect system may also be used to express a polypeptide of interest. For example, in one such system, Autographa californica nuclear polyhedrosis virus (AcNPV) is used as a vector to express foreign genes in *Spodoptera frugiperda* cells or in *Trichoplusia* larvae. The sequences encoding the polypeptide may be cloned into a non-essential region of the virus, such as the polyhedrin gene, and placed under control of the polyhedrin promoter. Successful insertion of the polypeptide-encoding sequence will render the polyhedrin gene inactive and produce recombinant virus lacking coat protein. The recombinant viruses may then be used to infect, for example, *S. frugiperda* cells or *Trichoplusia* larvae in which the polypeptide of interest may be expressed (Engelhard, E. K. *et al.* (1994) *Proc. Natl. Acad. Sci.* 91 :3224-3227).

In mammalian host cells, a number of viral-based expression systems are generally available. For example, in cases where an adenovirus is used as an expression vector, sequences encoding a polypeptide of interest may be ligated into an adenovirus transcription/translation complex consisting of the late promoter and tripartite leader sequence. Insertion in a non-essential E1 or E3 region of the viral genome may be used to obtain a viable virus which is capable of expressing the polypeptide in infected host cells (Logan, J. and Shenk, T. (1984) *Proc. Natl. Acad. Sci.* 81:3655-3659). In addition,

transcription enhancers, such as the Rous sarcoma virus (RSV) enhancer, may be used to increase expression in mammalian host cells.

Specific initiation signals may also be used to achieve more efficient translation of sequences encoding a polypeptide of interest. Such signals include the ATG initiation codon and adjacent sequences. In cases where sequences encoding the polypeptide, its initiation codon, and upstream sequences are inserted into the appropriate expression vector, no additional transcriptional or translational control signals may be needed. However, in cases where only coding sequence, or a portion thereof, is inserted, exogenous translational control signals including the ATG initiation codon should be provided. Furthermore, the initiation codon should be in the correct reading frame to ensure translation of the entire insert. Exogenous translational elements and initiation codons may be of various origins, both natural and synthetic. The efficiency of expression may be enhanced by the inclusion of enhancers which are appropriate for the particular cell system which is used, such as those described in the literature (Scharf, D. *et al.* (1994) *Results Probl. Cell Differ.* 20:125-162).

In addition, a host cell strain may be chosen for its ability to modulate the expression of the inserted sequences or to process the expressed protein in the desired fashion. Such modifications of the polypeptide include, but are not limited to, acetylation, carboxylation, glycosylation, phosphorylation, lipidation, and acylation. Post-translational processing which cleaves a "prepro" form of the protein may also be used to facilitate correct insertion, folding and/or function. Different host cells such as CHO, HeLa, MDCK, HEK293, and WI38, which have specific cellular machinery and characteristic mechanisms for such post-translational activities, may be chosen to ensure the correct modification and processing of the foreign protein.

For long-term, high-yield production of recombinant proteins, stable expression is generally preferred. For example, cell lines which stably express a polynucleotide of interest may be transformed using expression vectors which may contain viral origins of replication and/or endogenous expression elements and a selectable marker gene on the same or on a separate vector. Following the introduction of the vector, cells

may be allowed to grow for 1-2 days in an enriched media before they are switched to selective media. The purpose of the selectable marker is to confer resistance to selection, and its presence allows growth and recovery of cells which successfully express the introduced sequences. Resistant clones of stably transformed cells may be proliferated
5 using tissue culture techniques appropriate to the cell type.

Any number of selection systems may be used to recover transformed cell lines. These include, but are not limited to, the herpes simplex virus thymidine kinase (Wigler, M. *et al.* (1977) *Cell* 11:223-32) and adenine phosphoribosyltransferase (Lowy, I. *et al.* (1990) *Cell* 22:817-23) genes which can be employed in tk.sup.- or aprt.sup.- cells,
10 respectively. Also, antimetabolite, antibiotic or herbicide resistance can be used as the basis for selection; for example, dhfr which confers resistance to methotrexate (Wigler, M. *et al.* (1980) *Proc. Natl. Acad. Sci.* 77:3567-70); npt, which confers resistance to the aminoglycosides, neomycin and G-418 (Colbere-Garapin, F. *et al.* (1981) *J. Mol. Biol.* 150:1-14); and als or pat, which confer resistance to chlorsulfuron and phosphinotricin
15 acetyltransferase, respectively (Murry, *supra*). Additional selectable genes have been described, for example, trpB, which allows cells to utilize indole in place of tryptophan, or hisD, which allows cells to utilize histinol in place of histidine (Hartman, S. C. and R. C. Mulligan (1988) *Proc. Natl. Acad. Sci.* 85:8047-51). Recently, the use of visible markers has gained popularity with such markers as anthocyanins, beta-glucuronidase and its
20 substrate GUS, and luciferase and its substrate luciferin, being widely used not only to identify transformants, but also to quantify the amount of transient or stable protein expression attributable to a specific vector system (Rhodes, C. A. *et al.* (1995) *Methods Mol. Biol.* 55:121-131).

Although the presence/absence of marker gene expression suggests that the
25 gene of interest is also present, its presence and expression may need to be confirmed. For example, if the sequence encoding a polypeptide is inserted within a marker gene sequence, recombinant cells containing sequences can be identified by the absence of marker gene function. Alternatively, a marker gene can be placed in tandem with a polypeptide-encoding sequence under the control of a single promoter. Expression of the marker gene

in response to induction or selection usually indicates expression of the tandem gene as well.

Alternatively, host cells which contain and express a desired polynucleotide sequence may be identified by a variety of procedures known to those of skill in the art.

- 5 These procedures include, but are not limited to, DNA-DNA or DNA-RNA hybridizations and protein bioassay or immunoassay techniques which include membrane, solution, or chip based technologies for the detection and/or quantification of nucleic acid or protein.

A variety of protocols for detecting and measuring the expression of polynucleotide-encoded products, using either polyclonal or monoclonal antibodies
 10 specific for the product are known in the art. Examples include enzyme-linked immunosorbent assay (ELISA), radioimmunoassay (RIA), and fluorescence activated cell sorting (FACS). A two-site, monoclonal-based immunoassay utilizing monoclonal antibodies reactive to two non-interfering epitopes on a given polypeptide may be preferred for some applications, but a competitive binding assay may also be employed. These and
 15 other assays are described, among other places, in Hampton, R. *et al.* (1990; Serological Methods, a Laboratory Manual, APS Press, St Paul, Minn.) and Maddox, D. E. *et al.* (1983; *J. Exp. Med.* 158:1211-1216).

A wide variety of labels and conjugation techniques are known by those skilled in the art and may be used in various nucleic acid and amino acid assays. Means for
 20 producing labeled hybridization or PCR probes for detecting sequences related to polynucleotides include oligolabeling, nick translation, end-labeling or PCR amplification using a labeled nucleotide. Alternatively, the sequences, or any portions thereof may be cloned into a vector for the production of an mRNA probe. Such vectors are known in the art, are commercially available, and may be used to synthesize RNA probes in vitro by
 25 addition of an appropriate RNA polymerase such as T7, T3, or SP6 and labeled nucleotides. These procedures may be conducted using a variety of commercially available kits. Suitable reporter molecules or labels, which may be used include radionuclides, enzymes, fluorescent, chemiluminescent, or chromogenic agents as well as substrates, cofactors, inhibitors, magnetic particles, and the like.

Host cells transformed with a polynucleotide sequence of interest may be cultured under conditions suitable for the expression and recovery of the protein from cell culture. The protein produced by a recombinant cell may be secreted or contained intracellularly depending on the sequence and/or the vector used. As will be understood by

5 those of skill in the art, expression vectors containing polynucleotides of the invention may be designed to contain signal sequences which direct secretion of the encoded polypeptide through a prokaryotic or eukaryotic cell membrane. Other recombinant constructions may be used to join sequences encoding a polypeptide of interest to nucleotide sequence encoding a polypeptide domain which will facilitate purification of soluble proteins. Such

10 purification facilitating domains include, but are not limited to, metal chelating peptides such as histidine-tryptophan modules that allow purification on immobilized metals, protein A domains that allow purification on immobilized immunoglobulin, and the domain utilized in the FLAGS extension/affinity purification system (Immunex Corp., Seattle, Wash.). The inclusion of cleavable linker sequences such as those specific for Factor XA or

15 enterokinase (Invitrogen, San Diego, Calif.) between the purification domain and the encoded polypeptide may be used to facilitate purification. One such expression vector provides for expression of a fusion protein containing a polypeptide of interest and a nucleic acid encoding 6 histidine residues preceding a thioredoxin or an enterokinase cleavage site. The histidine residues facilitate purification on IMIAC (immobilized metal

20 ion affinity chromatography) as described in Porath, J. *et al.* (1992, *Prot. Exp. Purif.* 3:263-281) while the enterokinase cleavage site provides a means for purifying the desired polypeptide from the fusion protein. A discussion of vectors which contain fusion proteins is provided in Kroll, D. J. *et al.* (1993; *DNA Cell Biol.* 12:441-453).

In addition to recombinant production methods, polypeptides of the

25 invention, and fragments thereof, may be produced by direct peptide synthesis using solid-phase techniques (Merrifield J. (1963) *J. Am. Chem. Soc.* 85:2149-2154). Protein synthesis may be performed using manual techniques or by automation. Automated synthesis may be achieved, for example, using Applied Biosystems 431A Peptide Synthesizer (Perkin

Elmer). Alternatively, various fragments may be chemically synthesized separately and combined using chemical methods to produce the full length molecule.

SITE-SPECIFIC MUTAGENESIS

Site-specific mutagenesis is a technique useful in the preparation of individual peptides, or biologically functional equivalent polypeptides, through specific mutagenesis of the underlying polynucleotides that encode them. The technique, well-known to those of skill in the art, further provides a ready ability to prepare and test sequence variants, for example, incorporating one or more of the foregoing considerations, by introducing one or more nucleotide sequence changes into the DNA. Site-specific mutagenesis allows the production of mutants through the use of specific oligonucleotide sequences which encode the DNA sequence of the desired mutation, as well as a sufficient number of adjacent nucleotides, to provide a primer sequence of sufficient size and sequence complexity to form a stable duplex on both sides of the deletion junction being traversed. Mutations may be employed in a selected polynucleotide sequence to improve, alter, decrease, modify, or otherwise change the properties of the polynucleotide itself, and/or alter the properties, activity, composition, stability, or primary sequence of the encoded polypeptide.

In certain embodiments of the present invention, the inventors contemplate the mutagenesis of the disclosed polynucleotide sequences to alter one or more properties of the encoded polypeptide, such as the antigenicity of a polypeptide vaccine. The techniques of site-specific mutagenesis are well-known in the art, and are widely used to create variants of both polypeptides and polynucleotides. For example, site-specific mutagenesis is often used to alter a specific portion of a DNA molecule. In such embodiments, a primer comprising typically about 14 to about 25 nucleotides or so in length is employed, with about 5 to about 10 residues on both sides of the junction of the sequence being altered.

As will be appreciated by those of skill in the art, site-specific mutagenesis techniques have often employed a phage vector that exists in both a single stranded and

double stranded form. Typical vectors useful in site-directed mutagenesis include vectors such as the M13 phage. These phage are readily commercially-available and their use is generally well-known to those skilled in the art. Double-stranded plasmids are also routinely employed in site directed mutagenesis that eliminates the step of transferring the gene of interest from a plasmid to a phage.

In general, site-directed mutagenesis in accordance herewith is performed by first obtaining a single-stranded vector or melting apart of two strands of a double-stranded vector that includes within its sequence a DNA sequence that encodes the desired peptide. An oligonucleotide primer bearing the desired mutated sequence is prepared, generally synthetically. This primer is then annealed with the single-stranded vector, and subjected to DNA polymerizing enzymes such as *E. coli* polymerase I Klenow fragment, in order to complete the synthesis of the mutation-bearing strand. Thus, a heteroduplex is formed wherein one strand encodes the original non-mutated sequence and the second strand bears the desired mutation. This heteroduplex vector is then used to transform appropriate cells, such as *E. coli* cells, and clones are selected which include recombinant vectors bearing the mutated sequence arrangement.

The preparation of sequence variants of the selected peptide-encoding DNA segments using site-directed mutagenesis provides a means of producing potentially useful species and is not meant to be limiting as there are other ways in which sequence variants of peptides and the DNA sequences encoding them may be obtained. For example, recombinant vectors encoding the desired peptide sequence may be treated with mutagenic agents, such as hydroxylamine, to obtain sequence variants. Specific details regarding these methods and protocols are found in the teachings of Maloy *et al.*, 1994; Segal, 1976; Prokop and Bajpai, 1991; Kuby, 1994; and Maniatis *et al.*, 1982, each incorporated herein by reference, for that purpose.

As used herein, the term "oligonucleotide directed mutagenesis procedure" refers to template-dependent processes and vector-mediated propagation which result in an increase in the concentration of a specific nucleic acid molecule relative to its initial concentration, or in an increase in the concentration of a detectable signal, such as

amplification. As used herein, the term "oligonucleotide directed mutagenesis procedure" is intended to refer to a process that involves the template-dependent extension of a primer molecule. The term template dependent process refers to nucleic acid synthesis of an RNA or a DNA molecule wherein the sequence of the newly synthesized strand of nucleic acid is dictated by the well-known rules of complementary base pairing (see, for example, Watson, 1987). Typically, vector mediated methodologies involve the introduction of the nucleic acid fragment into a DNA or RNA vector, the clonal amplification of the vector, and the recovery of the amplified nucleic acid fragment. Examples of such methodologies are provided by U. S. Patent No. 4,237,224, specifically incorporated herein by reference in its entirety.

POLYNUCLEOTIDE AMPLIFICATION TECHNIQUES

A number of template dependent processes are available to amplify the target sequences of interest present in a sample. One of the best known amplification methods is the polymerase chain reaction (PCR™) which is described in detail in U.S. Patent Nos. 4,683,195, 4,683,202 and 4,800,159, each of which is incorporated herein by reference in its entirety. Briefly, in PCR™, two primer sequences are prepared which are complementary to regions on opposite complementary strands of the target sequence. An excess of deoxynucleoside triphosphates is added to a reaction mixture along with a DNA polymerase (e.g., *Taq* polymerase). If the target sequence is present in a sample, the primers will bind to the target and the polymerase will cause the primers to be extended along the target sequence by adding on nucleotides. By raising and lowering the temperature of the reaction mixture, the extended primers will dissociate from the target to form reaction products, excess primers will bind to the target and to the reaction product and the process is repeated. Preferably reverse transcription and PCR™ amplification procedure may be performed in order to quantify the amount of mRNA amplified. Polymerase chain reaction methodologies are well known in the art.

Another method for amplification is the ligase chain reaction (referred to as LCR), disclosed in Eur. Pat. Appl. Publ. No. 320,308 (specifically incorporated herein by

reference in its entirety). In LCR, two complementary probe pairs are prepared, and in the presence of the target sequence, each pair will bind to opposite complementary strands of the target such that they abut. In the presence of a ligase, the two probe pairs will link to form a single unit. By temperature cycling, as in PCR™, bound ligated units dissociate
 5 from the target and then serve as "target sequences" for ligation of excess probe pairs. U.S. Patent No. 4,883,750, incorporated herein by reference in its entirety, describes an alternative method of amplification similar to LCR for binding probe pairs to a target sequence.

Qbeta Replicase, described in PCT Intl. Pat. Appl. Publ. No.
 10 PCT/US87/00880, incorporated herein by reference in its entirety, may also be used as still another amplification method in the present invention. In this method, a replicative sequence of RNA that has a region complementary to that of a target is added to a sample in the presence of an RNA polymerase. The polymerase will copy the replicative sequence that can then be detected.

15 An isothermal amplification method, in which restriction endonucleases and ligases are used to achieve the amplification of target molecules that contain nucleotide 5'-[α-thio]triphosphates in one strand of a restriction site (Walker *et al.*, 1992, incorporated herein by reference in its entirety), may also be useful in the amplification of nucleic acids in the present invention.

20 Strand Displacement Amplification (SDA) is another method of carrying out isothermal amplification of nucleic acids which involves multiple rounds of strand displacement and synthesis, *i.e.* nick translation. A similar method, called Repair Chain Reaction (RCR) is another method of amplification which may be useful in the present invention and is involves annealing several probes throughout a region targeted for
 25 amplification, followed by a repair reaction in which only two of the four bases are present. The other two bases can be added as biotinylated derivatives for easy detection. A similar approach is used in SDA.

Sequences can also be detected using a cyclic probe reaction (CPR). In CPR, a probe having a 3' and 5' sequences of non-target DNA and an internal or "middle"

sequence of the target protein specific RNA is hybridized to DNA which is present in a sample. Upon hybridization, the reaction is treated with RNaseH, and the products of the probe are identified as distinctive products by generating a signal that is released after digestion. The original template is annealed to another cycling probe and the reaction is
 5 repeated. Thus, CPR involves amplifying a signal generated by hybridization of a probe to a target gene specific expressed nucleic acid.

Still other amplification methods described in Great Britain Pat. Appl. No. 2 202 328, and in PCT Intl. Pat. Appl. Publ. No. PCT/US89/01025, each of which is incorporated herein by reference in its entirety, may be used in accordance with the present
 10 invention. In the former application, "modified" primers are used in a PCR-like, template and enzyme dependent synthesis. The primers may be modified by labeling with a capture moiety (*e.g.*, biotin) and/or a detector moiety (*e.g.*, enzyme). In the latter application, an excess of labeled probes is added to a sample. In the presence of the target sequence, the probe binds and is cleaved catalytically. After cleavage, the target sequence is released
 15 intact to be bound by excess probe. Cleavage of the labeled probe signals the presence of the target sequence.

Other nucleic acid amplification procedures include transcription-based amplification systems (TAS) (Kwoh *et al.*, 1989; PCT Intl. Pat. Appl. Publ. No. WO 88/10315, incorporated herein by reference in its entirety), including nucleic acid sequence
 20 based amplification (NASBA) and 3SR. In NASBA, the nucleic acids can be prepared for amplification by standard phenol/chloroform extraction, heat denaturation of a sample, treatment with lysis buffer and minispin columns for isolation of DNA and RNA or guanidinium chloride extraction of RNA. These amplification techniques involve annealing a primer that has sequences specific to the target sequence. Following
 25 polymerization, DNA/RNA hybrids are digested with RNase H while double stranded DNA molecules are heat-denatured again. In either case the single stranded DNA is made fully double stranded by addition of second target-specific primer, followed by polymerization. The double stranded DNA molecules are then multiply transcribed by a polymerase such as T7 or SP6. In an isothermal cyclic reaction, the RNAs are reverse

transcribed into DNA, and transcribed once again with a polymerase such as T7 or SP6. The resulting products, whether truncated or complete, indicate target-specific sequences.

Eur. Pat. Appl. Publ. No. 329,822, incorporated herein by reference in its entirety, disclose a nucleic acid amplification process involving cyclically synthesizing
 5 single-stranded RNA ("ssRNA"), ssDNA, and double-stranded DNA (dsDNA), which may be used in accordance with the present invention. The ssRNA is a first template for a first primer oligonucleotide, which is elongated by reverse transcriptase (RNA-dependent DNA polymerase). The RNA is then removed from resulting DNA:RNA duplex by the action of ribonuclease H (RNase H, an RNase specific for RNA in a duplex with either DNA or
 10 RNA). The resultant ssDNA is a second template for a second primer, which also includes the sequences of an RNA polymerase promoter (exemplified by T7 RNA polymerase) 5' to its homology to its template. This primer is then extended by DNA polymerase (exemplified by the large "Klenow" fragment of *E. coli* DNA polymerase I), resulting as a double-stranded DNA ("dsDNA") molecule, having a sequence identical to that of the
 15 original RNA between the primers and having additionally, at one end, a promoter sequence. This promoter sequence can be used by the appropriate RNA polymerase to make many RNA copies of the DNA. These copies can then re-enter the cycle leading to very swift amplification. With proper choice of enzymes, this amplification can be done isothermally without addition of enzymes at each cycle. Because of the cyclical nature of
 20 this process, the starting sequence can be chosen to be in the form of either DNA or RNA.

PCT Intl. Pat. Appl. Publ. No. WO 89/06700, incorporated herein by reference in its entirety, disclose a nucleic acid sequence amplification scheme based on the hybridization of a promoter/primer sequence to a target single-stranded DNA ("ssDNA") followed by transcription of many RNA copies of the sequence. This scheme is not cyclic;
 25 *i.e.* new templates are not produced from the resultant RNA transcripts. Other amplification methods include "RACE" (Frohman, 1990), and "one-sided PCR" (Ohara, 1989) which are well-known to those of skill in the art.

Methods based on ligation of two (or more) oligonucleotides in the presence of nucleic acid having the sequence of the resulting "di-oligonucleotide", thereby

amplifying the di-oligonucleotide (Wu and Dean, 1996, incorporated herein by reference in its entirety), may also be used in the amplification of DNA sequences of the present invention.

BIOLOGICAL FUNCTIONAL EQUIVALENTS

5 Modification and changes may be made in the structure of the polynucleotides and polypeptides of the present invention and still obtain a functional molecule that encodes a polypeptide with desirable characteristics. As mentioned above, it is often desirable to introduce one or more mutations into a specific polynucleotide sequence. In certain circumstances, the resulting encoded polypeptide sequence is altered
10 by this mutation, or in other cases, the sequence of the polypeptide is unchanged by one or more mutations in the encoding polynucleotide.

When it is desirable to alter the amino acid sequence of a polypeptide to create an equivalent, or even an improved, second-generation molecule, the amino acid changes may be achieved by changing one or more of the codons of the encoding DNA
15 sequence, according to Table 1.

For example, certain amino acids may be substituted for other amino acids in a protein structure without appreciable loss of interactive binding capacity with structures such as, for example, antigen-binding regions of antibodies or binding sites on substrate molecules. Since it is the interactive capacity and nature of a protein that defines
20 that protein's biological functional activity, certain amino acid sequence substitutions can be made in a protein sequence, and, of course, its underlying DNA coding sequence, and nevertheless obtain a protein with like properties. It is thus contemplated by the inventors that various changes may be made in the peptide sequences of the disclosed compositions, or corresponding DNA sequences which encode said peptides without appreciable loss of
25 their biological utility or activity.

TABLE 1

Amino Acids			Codons						
Alanine	Ala	A	GCA	GCC	GCG	GCU			
Cysteine	Cys	C	UGC	UGU					
Aspartic acid	Asp	D	GAC	GAU					
Glutamic acid	Glu	E	GAA	GAG					
Phenylalanine	Phe	F	UUC	UUU					
Glycine	Gly	G	GGA	GGC	GGG	GGU			
Histidine	His	H	CAC	CAU					
Isoleucine	Ile	I	AUA	AUC	AUU				
Lysine	Lys	K	AAA	AAG					
Leucine	Leu	L	UUA	UUG	CUA	CUC	CUG	CUU	
Methionine	Met	M	AUG						
Asparagine	Asn	N	AAC	AAU					
Proline	Pro	P	CCA	CCC	CCG	CCU			
Glutamine	Gln	Q	CAA	CAG					
Arginine	Arg	R	AGA	AGG	CGA	CGC	CGG	CGU	
Serine	Ser	S	AGC	AGU	UCA	UCC	UCG	UCU	
Threonine	Thr	T	ACA	ACC	ACG	ACU			
Valine	Val	V	GUA	GUC	GUG	GUU			
Tryptophan	Trp	W	UGG						
Tyrosine	Tyr	Y	UAC	UAU					

In making such changes, the hydropathic index of amino acids may be considered. The importance of the hydropathic amino acid index in conferring interactive biologic function on a protein is generally understood in the art (Kyte and Doolittle, 1982, incorporated herein by reference). It is accepted that the relative hydropathic character of the amino acid contributes to the secondary structure of the resultant protein, which in turn defines the interaction of the protein with other molecules, for example, enzymes,

substrates, receptors, DNA, antibodies, antigens, and the like. Each amino acid has been assigned a hydropathic index on the basis of its hydrophobicity and charge characteristics (Kyte and Doolittle, 1982). These values are: isoleucine (+4.5); valine (+4.2); leucine (+3.8); phenylalanine (+2.8); cysteine/cystine (+2.5); methionine (+1.9); alanine (+1.8);
 5 glycine (−0.4); threonine (−0.7); serine (−0.8); tryptophan (−0.9); tyrosine (−1.3); proline (−1.6); histidine (−3.2); glutamate (−3.5); glutamine (−3.5); aspartate (−3.5); asparagine (−3.5); lysine (−3.9); and arginine (−4.5).

It is known in the art that certain amino acids may be substituted by other amino acids having a similar hydropathic index or score and still result in a protein with
 10 similar biological activity, *i.e.* still obtain a biological functionally equivalent protein. In making such changes, the substitution of amino acids whose hydropathic indices are within ± 2 is preferred, those within ± 1 are particularly preferred, and those within ± 0.5 are even more particularly preferred. It is also understood in the art that the substitution of like amino acids can be made effectively on the basis of hydrophilicity. U. S. Patent 4,554,101
 15 (specifically incorporated herein by reference in its entirety), states that the greatest local average hydrophilicity of a protein, as governed by the hydrophilicity of its adjacent amino acids, correlates with a biological property of the protein.

As detailed in U. S. Patent 4,554,101, the following hydrophilicity values have been assigned to amino acid residues: arginine (+3.0); lysine (+3.0); aspartate (+3.0 \pm 1); glutamate (+3.0 \pm 1); serine (+0.3); asparagine (+0.2); glutamine (+0.2); glycine (0);
 20 threonine (−0.4); proline (−0.5 \pm 1); alanine (−0.5); histidine (−0.5); cysteine (−1.0); methionine (−1.3); valine (−1.5); leucine (−1.8); isoleucine (−1.8); tyrosine (−2.3); phenylalanine (−2.5); tryptophan (−3.4). It is understood that an amino acid can be substituted for another having a similar hydrophilicity value and still obtain a biologically
 25 equivalent, and in particular, an immunologically equivalent protein. In such changes, the substitution of amino acids whose hydrophilicity values are within ± 2 is preferred, those within ± 1 are particularly preferred, and those within ± 0.5 are even more particularly preferred.

As outlined above, amino acid substitutions are generally therefore based on the relative similarity of the amino acid side-chain substituents, for example, their hydrophobicity, hydrophilicity, charge, size, and the like. Exemplary substitutions that take various of the foregoing characteristics into consideration are well known to those of skill in the art and include: arginine and lysine; glutamate and aspartate; serine and threonine; glutamine and asparagine; and valine, leucine and isoleucine.

In addition, any polynucleotide may be further modified to increase stability *in vivo*. Possible modifications include, but are not limited to, the addition of flanking sequences at the 5' and/or 3' ends; the use of phosphorothioate or 2' O-methyl rather than phosphodiesterase linkages in the backbone; and/or the inclusion of nontraditional bases such as inosine, queosine and wybutosine, as well as acetyl- methyl-, thio- and other modified forms of adenine, cytidine, guanine, thymine and uridine.

IN VIVO POLYNUCLEOTIDE DELIVERY TECHNIQUES

In additional embodiments, genetic constructs comprising one or more of the polynucleotides of the invention are introduced into cells *in vivo*. This may be achieved using any of a variety of well known approaches, several of which are outlined below for the purpose of illustration.

1. ADENOVIRUS

One of the preferred methods for *in vivo* delivery of one or more nucleic acid sequences involves the use of an adenovirus expression vector. "Adenovirus expression vector" is meant to include those constructs containing adenovirus sequences sufficient to (a) support packaging of the construct and (b) to express a polynucleotide that has been cloned therein in a sense or antisense orientation. Of course, in the context of an antisense construct, expression does not require that the gene product be synthesized.

The expression vector comprises a genetically engineered form of an adenovirus. Knowledge of the genetic organization of adenovirus, a 36 kb, linear, double-stranded DNA virus, allows substitution of large pieces of adenoviral DNA with foreign

sequences up to 7 kb (Grunhaus and Horwitz, 1992). In contrast to retrovirus, the adenoviral infection of host cells does not result in chromosomal integration because adenoviral DNA can replicate in an episomal manner without potential genotoxicity. Also, adenoviruses are structurally stable, and no genome rearrangement has been detected after
 5 extensive amplification. Adenovirus can infect virtually all epithelial cells regardless of their cell cycle stage. So far, adenoviral infection appears to be linked only to mild disease such as acute respiratory disease in humans.

Adenovirus is particularly suitable for use as a gene transfer vector because of its mid-sized genome, ease of manipulation, high titer, wide target-cell range and high
 10 infectivity. Both ends of the viral genome contain 100-200 base pair inverted repeats (ITRs), which are *cis* elements necessary for viral DNA replication and packaging. The early (E) and late (L) regions of the genome contain different transcription units that are divided by the onset of viral DNA replication. The E1 region (E1A and E1B) encodes proteins responsible for the regulation of transcription of the viral genome and a few
 15 cellular genes. The expression of the E2 region (E2A and E2B) results in the synthesis of the proteins for viral DNA replication. These proteins are involved in DNA replication, late gene expression and host cell shut-off (Renan, 1990). The products of the late genes, including the majority of the viral capsid proteins, are expressed only after significant processing of a single primary transcript issued by the major late promoter (MLP). The
 20 MLP, (located at 16.8 m.u.) is particularly efficient during the late phase of infection, and all the mRNA's issued from this promoter possess a 5'-tripartite leader (TPL) sequence which makes them preferred mRNA's for translation.

In a current system, recombinant adenovirus is generated from homologous recombination between shuttle vector and provirus vector. Due to the possible
 25 recombination between two proviral vectors, wild-type adenovirus may be generated from this process. Therefore, it is critical to isolate a single clone of virus from an individual plaque and examine its genomic structure.

Generation and propagation of the current adenovirus vectors, which are replication deficient, depend on a unique helper cell line, designated 293, which was

transformed from human embryonic kidney cells by Ad5 DNA fragments and constitutively expresses E1 proteins (Graham *et al.*, 1977). Since the E3 region is dispensable from the adenovirus genome (Jones and Shenk, 1978), the current adenovirus vectors, with the help of 293 cells, carry foreign DNA in either the E1, the D3 or both regions (Graham and Prevec, 1991). In nature, adenovirus can package approximately 105% of the wild-type genome (Ghosh-Choudhury *et al.*, 1987), providing capacity for about 2 extra kB of DNA. Combined with the approximately 5.5 kB of DNA that is replaceable in the E1 and E3 regions, the maximum capacity of the current adenovirus vector is under 7.5 kB, or about 15% of the total length of the vector. More than 80% of the adenovirus viral genome remains in the vector backbone and is the source of vector-borne cytotoxicity. Also, the replication deficiency of the E1-deleted virus is incomplete. For example, leakage of viral gene expression has been observed with the currently available vectors at high multiplicities of infection (MOI) (Mulligan, 1993).

Helper cell lines may be derived from human cells such as human embryonic kidney cells, muscle cells, hematopoietic cells or other human embryonic mesenchymal or epithelial cells. Alternatively, the helper cells may be derived from the cells of other mammalian species that are permissive for human adenovirus. Such cells include, *e.g.*, Vero cells or other monkey embryonic mesenchymal or epithelial cells. As stated above, the currently preferred helper cell line is 293.

Recently, Racher *et al.* (1995) disclosed improved methods for culturing 293 cells and propagating adenovirus. In one format, natural cell aggregates are grown by inoculating individual cells into 1 liter siliconized spinner flasks (Techne, Cambridge, UK) containing 100-200 ml of medium. Following stirring at 40 rpm, the cell viability is estimated with trypan blue. In another format, Fibra-Cel microcarriers (Bibby Sterlin, Stone, UK) (5 g/l) is employed as follows. A cell inoculum, resuspended in 5 ml of medium, is added to the carrier (50 ml) in a 250 ml Erlenmeyer flask and left stationary, with occasional agitation, for 1 to 4 h. The medium is then replaced with 50 ml of fresh medium and shaking initiated. For virus production, cells are allowed to grow to about 80% confluence, after which time the medium is replaced (to 25% of the final volume) and

adenovirus added at an MOI of 0.05. Cultures are left stationary overnight, following which the volume is increased to 100% and shaking commenced for another 72 h.

Other than the requirement that the adenovirus vector be replication defective, or at least conditionally defective, the nature of the adenovirus vector is not
 5 believed to be crucial to the successful practice of the invention. The adenovirus may be of any of the 42 different known serotypes or subgroups A-F. Adenovirus type 5 of subgroup C is the preferred starting material in order to obtain a conditional replication-defective adenovirus vector for use in the present invention, since Adenovirus type 5 is a human adenovirus about which a great deal of biochemical and genetic information is known, and
 10 it has historically been used for most constructions employing adenovirus as a vector.

As stated above, the typical vector according to the present invention is replication defective and will not have an adenovirus E1 region. Thus, it will be most convenient to introduce the polynucleotide encoding the gene of interest at the position from which the E1-coding sequences have been removed. However, the position of
 15 insertion of the construct within the adenovirus sequences is not critical to the invention. The polynucleotide encoding the gene of interest may also be inserted in lieu of the deleted E3 region in E3 replacement vectors as described by Karlsson *et al.* (1986) or in the E4 region where a helper cell line or helper virus complements the E4 defect.

Adenovirus is easy to grow and manipulate and exhibits broad host range *in vitro* and *in vivo*. This group of viruses can be obtained in high titers, *e.g.*, 10^9 - 10^{11} plaque-forming units per ml, and they are highly infective. The life cycle of adenovirus does not require integration into the host cell genome. The foreign genes delivered by adenovirus vectors are episomal and, therefore, have low genotoxicity to host cells. No side effects have been reported in studies of vaccination with wild-type adenovirus (Couch *et al.*, 1963;
 25 Top *et al.*, 1971), demonstrating their safety and therapeutic potential as *in vivo* gene transfer vectors.

Adenovirus vectors have been used in eukaryotic gene expression (Levrero *et al.*, 1991; Gomez-Foix *et al.*, 1992) and vaccine development (Grunhaus and Horwitz, 1992; Graham and Prevec, 1992). Recently, animal studies suggested that recombinant

adenovirus could be used for gene therapy (Stratford-Perricaudet and Perricaudet, 1991; Stratford-Perricaudet *et al.*, 1990; Rich *et al.*, 1993). Studies in administering recombinant adenovirus to different tissues include trachea instillation (Rosenfeld *et al.*, 1991; Rosenfeld *et al.*, 1992), muscle injection (Ragot *et al.*, 1993), peripheral intravenous
 5 injections (Herz and Gerard, 1993) and stereotactic inoculation into the brain (Le Gal La Salle *et al.*, 1993).

2. RETROVIRUSES

The retroviruses are a group of single-stranded RNA viruses characterized by an ability to convert their RNA to double-stranded DNA in infected cells by a process of
 10 reverse-transcription (Coffin, 1990). The resulting DNA then stably integrates into cellular chromosomes as a provirus and directs synthesis of viral proteins. The integration results in the retention of the viral gene sequences in the recipient cell and its descendants. The retroviral genome contains three genes, gag, pol, and env that code for capsid proteins, polymerase enzyme, and envelope components, respectively. A sequence found upstream
 15 from the gag gene contains a signal for packaging of the genome into virions. Two long terminal repeat (LTR) sequences are present at the 5' and 3' ends of the viral genome. These contain strong promoter and enhancer sequences and are also required for integration in the host cell genome (Coffin, 1990).

In order to construct a retroviral vector, a nucleic acid encoding one or more
 20 oligonucleotide or polynucleotide sequences of interest is inserted into the viral genome in the place of certain viral sequences to produce a virus that is replication-defective. In order to produce virions, a packaging cell line containing the gag, pol, and env genes but without the LTR and packaging components is constructed (Mann *et al.*, 1983). When a recombinant plasmid containing a cDNA, together with the retroviral LTR and packaging
 25 sequences is introduced into this cell line (by calcium phosphate precipitation for example), the packaging sequence allows the RNA transcript of the recombinant plasmid to be packaged into viral particles, which are then secreted into the culture media (Nicolas and Rubenstein, 1988; Temin, 1986; Mann *et al.*, 1983). The media containing the

recombinant retroviruses is then collected, optionally concentrated, and used for gene transfer. Retroviral vectors are able to infect a broad variety of cell types. However, integration and stable expression require the division of host cells (Paskind *et al.*, 1975).

A novel approach designed to allow specific targeting of retrovirus vectors was recently developed based on the chemical modification of a retrovirus by the chemical addition of lactose residues to the viral envelope. This modification could permit the specific infection of hepatocytes *via* sialoglycoprotein receptors.

A different approach to targeting of recombinant retroviruses was designed in which biotinylated antibodies against a retroviral envelope protein and against a specific cell receptor were used. The antibodies were coupled *via* the biotin components by using streptavidin (Roux *et al.*, 1989). Using antibodies against major histocompatibility complex class I and class II antigens, they demonstrated the infection of a variety of human cells that bore those surface antigens with an ecotropic virus *in vitro* (Roux *et al.*, 1989).

3. ADENO-ASSOCIATED VIRUSES

AAV (Ridgeway, 1988; Hermonat and Muzyczka, 1984) is a parovirus, discovered as a contamination of adenoviral stocks. It is a ubiquitous virus (antibodies are present in 85% of the US human population) that has not been linked to any disease. It is also classified as a dependovirus, because its replications is dependent on the presence of a helper virus, such as adenovirus. Five serotypes have been isolated, of which AAV-2 is the best characterized. AAV has a single-stranded linear DNA that is encapsidated into capsid proteins VP1, VP2 and VP3 to form an icosahedral virion of 20 to 24 nm in diameter (Muzyczka and McLaughlin, 1988).

The AAV DNA is approximately 4.7 kilobases long. It contains two open reading frames and is flanked by two ITRs. There are two major genes in the AAV genome: *rep* and *cap*. The *rep* gene codes for proteins responsible for viral replications, whereas *cap* codes for capsid protein VP1-3. Each ITR forms a T-shaped hairpin structure. These terminal repeats are the only essential *cis* components of the AAV for chromosomal integration. Therefore, the AAV can be used as a vector with all viral coding

sequences removed and replaced by the cassette of genes for delivery. Three viral promoters have been identified and named p5, p19, and p40, according to their map position. Transcription from p5 and p19 results in production of rep proteins, and transcription from p40 produces the capsid proteins (Hermonat and Muzyczka, 1984).

5 There are several factors that prompted researchers to study the possibility of using rAAV as an expression vector. One is that the requirements for delivering a gene to integrate into the host chromosome are surprisingly few. It is necessary to have the 145-bp ITRs, which are only 6% of the AAV genome. This leaves room in the vector to assemble a 4.5-kb DNA insertion. While this carrying capacity may prevent the AAV from
10 delivering large genes, it is amply suited for delivering the antisense constructs of the present invention.

AAV is also a good choice of delivery vehicles due to its safety. There is a relatively complicated rescue mechanism: not only wild type adenovirus but also AAV genes are required to mobilize rAAV. Likewise, AAV is not pathogenic and not associated
15 with any disease. The removal of viral coding sequences minimizes immune reactions to viral gene expression, and therefore, rAAV does not evoke an inflammatory response.

4. OTHER VIRAL VECTORS AS EXPRESSION CONSTRUCTS

Other viral vectors may be employed as expression constructs in the present invention for the delivery of oligonucleotide or polynucleotide sequences to a host cell.
20 Vectors derived from viruses such as vaccinia virus (Ridgeway, 1988; Coupar *et al.*, 1988), lentiviruses, polio viruses and herpes viruses may be employed. They offer several attractive features for various mammalian cells (Friedmann, 1989; Ridgeway, 1988; Coupar *et al.*, 1988; Horwich *et al.*, 1990).

With the recent recognition of defective hepatitis B viruses, new insight was
25 gained into the structure-function relationship of different viral sequences. *In vitro* studies showed that the virus could retain the ability for helper-dependent packaging and reverse transcription despite the deletion of up to 80% of its genome (Horwich *et al.*, 1990). This suggested that large portions of the genome could be replaced with foreign genetic

material. The hepatotropism and persistence (integration) were particularly attractive properties for liver-directed gene transfer. Chang *et al.* (1991) introduced the chloramphenicol acetyltransferase (CAT) gene into duck hepatitis B virus genome in the place of the polymerase, surface, and pre-surface coding sequences. It was cotransfected with wild-type virus into an avian hepatoma cell line. Culture media containing high titers of the recombinant virus were used to infect primary duckling hepatocytes. Stable CAT gene expression was detected for at least 24 days after transfection (Chang *et al.*, 1991).

5. NON-VIRAL VECTORS

In order to effect expression of the oligonucleotide or polynucleotide sequences of the present invention, the expression construct must be delivered into a cell. This delivery may be accomplished *in vitro*, as in laboratory procedures for transforming cells lines, or *in vivo* or *ex vivo*, as in the treatment of certain disease states. As described above, one preferred mechanism for delivery is *via* viral infection where the expression construct is encapsulated in an infectious viral particle.

Once the expression construct has been delivered into the cell the nucleic acid encoding the desired oligonucleotide or polynucleotide sequences may be positioned and expressed at different sites. In certain embodiments, the nucleic acid encoding the construct may be stably integrated into the genome of the cell. This integration may be in the specific location and orientation *via* homologous recombination (gene replacement) or it may be integrated in a random, non-specific location (gene augmentation). In yet further embodiments, the nucleic acid may be stably maintained in the cell as a separate, episomal segment of DNA. Such nucleic acid segments or "episomes" encode sequences sufficient to permit maintenance and replication independent of or in synchronization with the host cell cycle. How the expression construct is delivered to a cell and where in the cell the nucleic acid remains is dependent on the type of expression construct employed.

In certain embodiments of the invention, the expression construct comprising one or more oligonucleotide or polynucleotide sequences may simply consist of naked recombinant DNA or plasmids. Transfer of the construct may be performed by any

of the methods mentioned above which physically or chemically permeabilize the cell membrane. This is particularly applicable for transfer *in vitro* but it may be applied to *in vivo* use as well. Dubensky *et al.* (1984) successfully injected polyomavirus DNA in the form of calcium phosphate precipitates into liver and spleen of adult and newborn mice demonstrating active viral replication and acute infection. Benvenisty and Reshef (1986) also demonstrated that direct intraperitoneal injection of calcium phosphate-precipitated plasmids results in expression of the transfected genes. It is envisioned that DNA encoding a gene of interest may also be transferred in a similar manner *in vivo* and express the gene product.

Another embodiment of the invention for transferring a naked DNA expression construct into cells may involve particle bombardment. This method depends on the ability to accelerate DNA-coated microprojectiles to a high velocity allowing them to pierce cell membranes and enter cells without killing them (Klein *et al.*, 1987). Several devices for accelerating small particles have been developed. One such device relies on a high voltage discharge to generate an electrical current, which in turn provides the motive force (Yang *et al.*, 1990). The microprojectiles used have consisted of biologically inert substances such as tungsten or gold beads.

Selected organs including the liver, skin, and muscle tissue of rats and mice have been bombarded *in vivo* (Yang *et al.*, 1990; Zelenin *et al.*, 1991). This may require surgical exposure of the tissue or cells, to eliminate any intervening tissue between the gun and the target organ, *i.e. ex vivo* treatment. Again, DNA encoding a particular gene may be delivered *via* this method and still be incorporated by the present invention.

ANTISENSE OLIGONUCLEOTIDES

The end result of the flow of genetic information is the synthesis of protein. DNA is transcribed by polymerases into messenger RNA and translated on the ribosome to yield a folded, functional protein. Thus there are several steps along the route where protein synthesis can be inhibited. The native DNA segment coding for a polypeptide described herein, as all such mammalian DNA strands, has two strands: a sense strand and

an antisense strand held together by hydrogen bonding. The messenger RNA coding for polypeptide has the same nucleotide sequence as the sense DNA strand except that the DNA thymidine is replaced by uridine. Thus, synthetic antisense nucleotide sequences will bind to a mRNA and inhibit expression of the protein encoded by that mRNA.

5 The targeting of antisense oligonucleotides to mRNA is thus one mechanism to shut down protein synthesis, and, consequently, represents a powerful and targeted therapeutic approach. For example, the synthesis of polygalacturonase and the muscarine type 2 acetylcholine receptor are inhibited by antisense oligonucleotides directed to their respective mRNA sequences (U. S. Patent 5,739,119 and U. S. Patent 5,759,829, each
10 specifically incorporated herein by reference in its entirety). Further, examples of antisense inhibition have been demonstrated with the nuclear protein cyclin, the multiple drug resistance gene (MDG1), ICAM-1, E-selectin, STK-1, striatal GABA_A receptor and human EGF (Jaskulski *et al.*, 1988; Vasanthakumar and Ahmed, 1989; Peris *et al.*, 1998; U. S. Patent 5,801,154; U. S. Patent 5,789,573; U. S. Patent 5,718,709 and U. S. Patent
15 5,610,288, each specifically incorporated herein by reference in its entirety). Antisense constructs have also been described that inhibit and can be used to treat a variety of abnormal cellular proliferations, *e.g.* cancer (U. S. Patent 5,747,470; U. S. Patent 5,591,317 and U. S. Patent 5,783,683, each specifically incorporated herein by reference in its entirety).

20 Therefore, in exemplary embodiments, the invention provides oligonucleotide sequences that comprise all, or a portion of, any sequence that is capable of specifically binding to polynucleotide sequence described herein, or a complement thereof. In one embodiment, the antisense oligonucleotides comprise DNA or derivatives thereof. In another embodiment, the oligonucleotides comprise RNA or derivatives thereof. In a
25 third embodiment, the oligonucleotides are modified DNAs comprising a phosphorothioated modified backbone. In a fourth embodiment, the oligonucleotide sequences comprise peptide nucleic acids or derivatives thereof. In each case, preferred compositions comprise a sequence region that is complementary, and more preferably

substantially-complementary, and even more preferably, completely complementary to one or more portions of polynucleotides disclosed herein.

Selection of antisense compositions specific for a given gene sequence is based upon analysis of the chosen target sequence (*i.e.* in these illustrative examples the rat
5 and human sequences) and determination of secondary structure, T_m , binding energy, relative stability, and antisense compositions were selected based upon their relative inability to form dimers, hairpins, or other secondary structures that would reduce or prohibit specific binding to the target mRNA in a host cell.

Highly preferred target regions of the mRNA, are those which are at or near
10 the AUG translation initiation codon, and those sequences which were substantially complementary to 5' regions of the mRNA. These secondary structure analyses and target site selection considerations were performed using v.4 of the OLIGO primer analysis software (Rychlik, 1997) and the BLASTN 2.0.5 algorithm software (Altschul *et al.*, 1997).

The use of an antisense delivery method employing a short peptide vector,
15 termed MPG (27 residues), is also contemplated. The MPG peptide contains a hydrophobic domain derived from the fusion sequence of HIV gp41 and a hydrophilic domain from the nuclear localization sequence of SV40 T-antigen (Morris *et al.*, 1997). It has been demonstrated that several molecules of the MPG peptide coat the antisense oligonucleotides and can be delivered into cultured mammalian cells in less than 1 hour
20 with relatively high efficiency (90%). Further, the interaction with MPG strongly increases both the stability of the oligonucleotide to nuclease and the ability to cross the plasma membrane (Morris *et al.*, 1997).

RIBOZYMES

Although proteins traditionally have been used for catalysis of nucleic acids,
25 another class of macromolecules has emerged as useful in this endeavor. Ribozymes are RNA-protein complexes that cleave nucleic acids in a site-specific fashion. Ribozymes have specific catalytic domains that possess endonuclease activity (Kim and Cech, 1987; Gerlach *et al.*, 1987; Forster and Symons, 1987). For example, a large number of

ribozymes accelerate phosphoester transfer reactions with a high degree of specificity, often cleaving only one of several phosphoesters in an oligonucleotide substrate (Cech *et al.*, 1981; Michel and Westhof, 1990; Reinhold-Hurek and Shub, 1992). This specificity has been attributed to the requirement that the substrate bind via specific base-pairing
 5 interactions to the internal guide sequence ("IGS") of the ribozyme prior to chemical reaction.

Ribozyme catalysis has primarily been observed as part of sequence-specific cleavage/ligation reactions involving nucleic acids (Joyce, 1989; Cech *et al.*, 1981). For example, U. S. Patent No. 5,354,855 (specifically incorporated herein by reference) reports
 10 that certain ribozymes can act as endonucleases with a sequence specificity greater than that of known ribonucleases and approaching that of the DNA restriction enzymes. Thus, sequence-specific ribozyme-mediated inhibition of gene expression may be particularly suited to therapeutic applications (Scanlon *et al.*, 1991; Sarver *et al.*, 1990). Recently, it was reported that ribozymes elicited genetic changes in some cells lines to which they were
 15 applied; the altered genes included the oncogenes *H-ras*, *c-fos* and genes of HIV. Most of this work involved the modification of a target mRNA, based on a specific mutant codon that is cleaved by a specific ribozyme.

Six basic varieties of naturally-occurring enzymatic RNAs are known presently. Each can catalyze the hydrolysis of RNA phosphodiester bonds *in trans* (and
 20 thus can cleave other RNA molecules) under physiological conditions. In general, enzymatic nucleic acids act by first binding to a target RNA. Such binding occurs through the target binding portion of a enzymatic nucleic acid which is held in close proximity to an enzymatic portion of the molecule that acts to cleave the target RNA. Thus, the enzymatic nucleic acid first recognizes and then binds a target RNA through complementary base-
 25 pairing, and once bound to the correct site, acts enzymatically to cut the target RNA. Strategic cleavage of such a target RNA will destroy its ability to direct synthesis of an encoded protein. After an enzymatic nucleic acid has bound and cleaved its RNA target, it is released from that RNA to search for another target and can repeatedly bind and cleave new targets.

The enzymatic nature of a ribozyme is advantageous over many technologies, such as antisense technology (where a nucleic acid molecule simply binds to a nucleic acid target to block its translation) since the concentration of ribozyme necessary to affect a therapeutic treatment is lower than that of an antisense oligonucleotide. This advantage reflects the ability of the ribozyme to act enzymatically. Thus, a single ribozyme molecule is able to cleave many molecules of target RNA. In addition, the ribozyme is a highly specific inhibitor, with the specificity of inhibition depending not only on the base pairing mechanism of binding to the target RNA, but also on the mechanism of target RNA cleavage. Single mismatches, or base-substitutions, near the site of cleavage can completely eliminate catalytic activity of a ribozyme. Similar mismatches in antisense molecules do not prevent their action (Woolf *et al.*, 1992). Thus, the specificity of action of a ribozyme is greater than that of an antisense oligonucleotide binding the same RNA site.

The enzymatic nucleic acid molecule may be formed in a hammerhead, hairpin, a hepatitis δ virus, group I intron or RNaseP RNA (in association with an RNA guide sequence) or Neurospora VS RNA motif. Examples of hammerhead motifs are described by Rossi *et al.* (1992). Examples of hairpin motifs are described by Hampel *et al.* (Eur. Pat. Appl. Publ. No. EP 0360257), Hampel and Tritz (1989), Hampel *et al.* (1990) and U. S. Patent 5,631,359 (specifically incorporated herein by reference). An example of the hepatitis δ virus motif is described by Perrotta and Been (1992); an example of the RNaseP motif is described by Guerrier-Takada *et al.* (1983); Neurospora VS RNA ribozyme motif is described by Collins (Saville and Collins, 1990; Saville and Collins, 1991; Collins and Olive, 1993); and an example of the Group I intron is described in (U. S. Patent 4,987,071, specifically incorporated herein by reference). All that is important in an enzymatic nucleic acid molecule of this invention is that it has a specific substrate binding site which is complementary to one or more of the target gene RNA regions, and that it have nucleotide sequences within or surrounding that substrate binding site which impart an RNA cleaving activity to the molecule. Thus the ribozyme constructs need not be limited to specific motifs mentioned herein.

In certain embodiments, it may be important to produce enzymatic cleaving agents which exhibit a high degree of specificity for the RNA of a desired target, such as one of the sequences disclosed herein. The enzymatic nucleic acid molecule is preferably targeted to a highly conserved sequence region of a target mRNA. Such enzymatic nucleic acid molecules can be delivered exogenously to specific cells as required. Alternatively, the ribozymes can be expressed from DNA or RNA vectors that are delivered to specific cells.

Small enzymatic nucleic acid motifs (e.g., of the hammerhead or the hairpin structure) may also be used for exogenous delivery. The simple structure of these molecules increases the ability of the enzymatic nucleic acid to invade targeted regions of the mRNA structure. Alternatively, catalytic RNA molecules can be expressed within cells from eukaryotic promoters (e.g., Scanlon *et al.*, 1991; Kashani-Sabet *et al.*, 1992; Dropulic *et al.*, 1992; Weerasinghe *et al.*, 1991; Ojwang *et al.*, 1992; Chen *et al.*, 1992; Sarver *et al.*, 1990). Those skilled in the art realize that any ribozyme can be expressed in eukaryotic cells from the appropriate DNA vector. The activity of such ribozymes can be augmented by their release from the primary transcript by a second ribozyme (Int. Pat. Appl. Publ. No. WO 93/23569, and Int. Pat. Appl. Publ. No. WO 94/02595, both hereby incorporated by reference; Ohkawa *et al.*, 1992; Taira *et al.*, 1991; and Ventura *et al.*, 1993).

Ribozymes may be added directly, or can be complexed with cationic lipids, lipid complexes, packaged within liposomes, or otherwise delivered to target cells. The RNA or RNA complexes can be locally administered to relevant tissues *ex vivo*, or *in vivo* through injection, aerosol inhalation, infusion pump or stent, with or without their incorporation in biopolymers.

Ribozymes may be designed as described in Int. Pat. Appl. Publ. No. WO 93/23569 and Int. Pat. Appl. Publ. No. WO 94/02595, each specifically incorporated herein by reference) and synthesized to be tested *in vitro* and *in vivo*, as described. Such ribozymes can also be optimized for delivery. While specific examples are provided, those in the art will recognize that equivalent RNA targets in other species can be utilized when necessary.

Hammerhead or hairpin ribozymes may be individually analyzed by computer folding (Jaeger *et al.*, 1989) to assess whether the ribozyme sequences fold into the appropriate secondary structure. Those ribozymes with unfavorable intramolecular interactions between the binding arms and the catalytic core are eliminated from
 5 consideration. Varying binding arm lengths can be chosen to optimize activity. Generally, at least 5 or so bases on each arm are able to bind to, or otherwise interact with, the target RNA.

Ribozymes of the hammerhead or hairpin motif may be designed to anneal to various sites in the mRNA message, and can be chemically synthesized. The method of
 10 synthesis used follows the procedure for normal RNA synthesis as described in Usman *et al.* (1987) and in Scaringe *et al.* (1990) and makes use of common nucleic acid protecting and coupling groups, such as dimethoxytrityl at the 5'-end, and phosphoramidites at the 3'-end. Average stepwise coupling yields are typically >98%. Hairpin ribozymes may be synthesized in two parts and annealed to reconstruct an active
 15 ribozyme (Chowrira and Burke, 1992). Ribozymes may be modified extensively to enhance stability by modification with nuclease resistant groups, for example, 2'-amino, 2'-C-allyl, 2'-fluoro, 2'-O-methyl, 2'-H (for a review see *e.g.*, Usman and Cedergren, 1992). Ribozymes may be purified by gel electrophoresis using general methods or by high pressure liquid chromatography and resuspended in water.

20 Ribozyme activity can be optimized by altering the length of the ribozyme binding arms, or chemically synthesizing ribozymes with modifications that prevent their degradation by serum ribonucleases (see *e.g.*, Int. Pat. Appl. Publ. No. WO 92/07065; Perrault *et al.*, 1990; Pieken *et al.*, 1991; Usman and Cedergren, 1992; Int. Pat. Appl. Publ. No. WO 93/15187; Int. Pat. Appl. Publ. No. WO 91/03162; Eur. Pat. Appl. Publ.
 25 No. 92110298.4; U. S. Patent 5,334,711; and Int. Pat. Appl. Publ. No. WO 94/13688, which describe various chemical modifications that can be made to the sugar moieties of enzymatic RNA molecules), modifications which enhance their efficacy in cells, and removal of stem II bases to shorten RNA synthesis times and reduce chemical requirements.

Sullivan *et al.* (Int. Pat. Appl. Publ. No. WO 94/02595) describes the general methods for delivery of enzymatic RNA molecules. Ribozymes may be administered to cells by a variety of methods known to those familiar to the art, including, but not restricted to, encapsulation in liposomes, by iontophoresis, or by incorporation into
 5 other vehicles, such as hydrogels, cyclodextrins, biodegradable nanocapsules, and bioadhesive microspheres. For some indications, ribozymes may be directly delivered *ex vivo* to cells or tissues with or without the aforementioned vehicles. Alternatively, the RNA/vehicle combination may be locally delivered by direct inhalation, by direct injection or by use of a catheter, infusion pump or stent. Other routes of delivery include, but are not
 10 limited to, intravascular, intramuscular, subcutaneous or joint injection, aerosol inhalation, oral (tablet or pill form), topical, systemic, ocular, intraperitoneal and/or intrathecal delivery. More detailed descriptions of ribozyme delivery and administration are provided in Int. Pat. Appl. Publ. No. WO 94/02595 and Int. Pat. Appl. Publ. No. WO 93/23569, each specifically incorporated herein by reference.

15 Another means of accumulating high concentrations of a ribozyme(s) within cells is to incorporate the ribozyme-encoding sequences into a DNA expression vector. Transcription of the ribozyme sequences are driven from a promoter for eukaryotic RNA polymerase I (pol I), RNA polymerase II (pol II), or RNA polymerase III (pol III). Transcripts from pol II or pol III promoters will be expressed at high levels in all cells; the
 20 levels of a given pol II promoter in a given cell type will depend on the nature of the gene regulatory sequences (enhancers, silencers, *etc.*) present nearby. Prokaryotic RNA polymerase promoters may also be used, providing that the prokaryotic RNA polymerase enzyme is expressed in the appropriate cells (Elroy-Stein and Moss, 1990; Gao and Huang, 1993; Lieber *et al.*, 1993; Zhou *et al.*, 1990). Ribozymes expressed from such promoters
 25 can function in mammalian cells (*e.g.* Kashani-Saber *et al.*, 1992; Ojwang *et al.*, 1992; Chen *et al.*, 1992; Yu *et al.*, 1993; L'Huillier *et al.*, 1992; Lisiewicz *et al.*, 1993). Such transcription units can be incorporated into a variety of vectors for introduction into mammalian cells, including but not restricted to, plasmid DNA vectors, viral DNA vectors

(such as adenovirus or adeno-associated vectors), or viral RNA vectors (such as retroviral, semliki forest virus, sindbis virus vectors).

Ribozymes may be used as diagnostic tools to examine genetic drift and mutations within diseased cells. They can also be used to assess levels of the target RNA molecule. The close relationship between ribozyme activity and the structure of the target RNA allows the detection of mutations in any region of the molecule which alters the base-pairing and three-dimensional structure of the target RNA. By using multiple ribozymes, one may map nucleotide changes which are important to RNA structure and function *in vitro*, as well as in cells and tissues. Cleavage of target RNAs with ribozymes may be used to inhibit gene expression and define the role (essentially) of specified gene products in the progression of disease. In this manner, other genetic targets may be defined as important mediators of the disease. These studies will lead to better treatment of the disease progression by affording the possibility of combinational therapies (*e.g.*, multiple ribozymes targeted to different genes, ribozymes coupled with known small molecule inhibitors, or intermittent treatment with combinations of ribozymes and/or other chemical or biological molecules). Other *in vitro* uses of ribozymes are well known in the art, and include detection of the presence of mRNA associated with an IL-5 related condition. Such RNA is detected by determining the presence of a cleavage product after treatment with a ribozyme using standard methodology.

20 PEPTIDE NUCLEIC ACIDS

In certain embodiments, the inventors contemplate the use of peptide nucleic acids (PNAs) in the practice of the methods of the invention. PNA is a DNA mimic in which the nucleobases are attached to a pseudopeptide backbone (Good and Nielsen, 1997). PNA is able to be utilized in a number methods that traditionally have used RNA or DNA. Often PNA sequences perform better in techniques than the corresponding RNA or DNA sequences and have utilities that are not inherent to RNA or DNA. A review of PNA including methods of making, characteristics of, and methods of using, is provided by Corey (1997) and is incorporated herein by reference. As such, in certain embodiments,

one may prepare PNA sequences that are complementary to one or more portions of the ACE mRNA sequence, and such PNA compositions may be used to regulate, alter, decrease, or reduce the translation of ACE-specific mRNA, and thereby alter the level of ACE activity in a host cell to which such PNA compositions have been administered.

5 PNAs have 2-aminoethyl-glycine linkages replacing the normal phosphodiester backbone of DNA (Nielsen *et al.*, 1991; Hanvey *et al.*, 1992; Hyrup and Nielsen, 1996; Neilsen, 1996). This chemistry has three important consequences: firstly, in contrast to DNA or phosphorothioate oligonucleotides, PNAs are neutral molecules; secondly, PNAs are achiral, which avoids the need to develop a stereoselective synthesis; 10 and thirdly, PNA synthesis uses standard Boc (Dueholm *et al.*, 1994) or Fmoc (Thomson *et al.*, 1995) protocols for solid-phase peptide synthesis, although other methods, including a modified Merrifield method, have been used (Christensen *et al.*, 1995).

PNA monomers or ready-made oligomers are commercially available from PerSeptive Biosystems (Framingham, MA). PNA syntheses by either Boc or Fmoc 15 protocols are straightforward using manual or automated protocols (Norton *et al.*, 1995). The manual protocol lends itself to the production of chemically modified PNAs or the simultaneous synthesis of families of closely related PNAs.

As with peptide synthesis, the success of a particular PNA synthesis will depend on the properties of the chosen sequence. For example, while in theory PNAs can 20 incorporate any combination of nucleotide bases, the presence of adjacent purines can lead to deletions of one or more residues in the product. In expectation of this difficulty, it is suggested that, in producing PNAs with adjacent purines, one should repeat the coupling of residues likely to be added inefficiently. This should be followed by the purification of PNAs by reverse-phase high-pressure liquid chromatography (Norton *et al.*, 1995) 25 providing yields and purity of product similar to those observed during the synthesis of peptides.

Modifications of PNAs for a given application may be accomplished by coupling amino acids during solid-phase synthesis or by attaching compounds that contain a carboxylic acid group to the exposed N-terminal amine. Alternatively, PNAs can be

modified after synthesis by coupling to an introduced lysine or cysteine. The ease with which PNAs can be modified facilitates optimization for better solubility or for specific functional requirements. Once synthesized, the identity of PNAs and their derivatives can be confirmed by mass spectrometry. Several studies have made and utilized modifications of PNAs (Norton *et al.*, 1995; Haaime *et al.*, 1996; Stetsenko *et al.*, 1996; Petersen *et al.*, 1995; Ulmann *et al.*, 1996; Koch *et al.*, 1995; Orum *et al.*, 1995; Footer *et al.*, 1996; Griffith *et al.*, 1995; Kremsky *et al.*, 1996; Pardridge *et al.*, 1995; Boffa *et al.*, 1995; Landsdorp *et al.*, 1996; Gambacorti-Passerini *et al.*, 1996; Armitage *et al.*, 1997; Seeger *et al.*, 1997; Ruskowski *et al.*, 1997). U.S. Patent No. 5,700,922 discusses PNA-DNA-PNA chimeric molecules and their uses in diagnostics, modulating protein in organisms, and treatment of conditions susceptible to therapeutics.

In contrast to DNA and RNA, which contain negatively charged linkages, the PNA backbone is neutral. In spite of this dramatic alteration, PNAs recognize complementary DNA and RNA by Watson-Crick pairing (Egholm *et al.*, 1993), validating the initial modeling by Nielsen *et al.* (1991). PNAs lack 3' to 5' polarity and can bind in either parallel or antiparallel fashion, with the antiparallel mode being preferred (Egholm *et al.*, 1993).

Hybridization of DNA oligonucleotides to DNA and RNA is destabilized by electrostatic repulsion between the negatively charged phosphate backbones of the complementary strands. By contrast, the absence of charge repulsion in PNA-DNA or PNA-RNA duplexes increases the melting temperature (T_m) and reduces the dependence of T_m on the concentration of mono- or divalent cations (Nielsen *et al.*, 1991). The enhanced rate and affinity of hybridization are significant because they are responsible for the surprising ability of PNAs to perform strand invasion of complementary sequences within relaxed double-stranded DNA. In addition, the efficient hybridization at inverted repeats suggests that PNAs can recognize secondary structure effectively within double-stranded DNA. Enhanced recognition also occurs with PNAs immobilized on surfaces, and Wang *et al.* have shown that support-bound PNAs can be used to detect hybridization events (Wang *et al.*, 1996).

One might expect that tight binding of PNAs to complementary sequences would also increase binding to similar (but not identical) sequences, reducing the sequence specificity of PNA recognition. As with DNA hybridization, however, selective recognition can be achieved by balancing oligomer length and incubation temperature.

5 Moreover, selective hybridization of PNAs is encouraged by PNA-DNA hybridization being less tolerant of base mismatches than DNA-DNA hybridization. For example, a single mismatch within a 16 bp PNA-DNA duplex can reduce the T_m by up to 15°C (Egholm *et al.*, 1993). This high level of discrimination has allowed the development of several PNA-based strategies for the analysis of point mutations (Wang *et al.*, 1996;

10 Carlsson *et al.*, 1996; Thiede *et al.*, 1996; Webb and Hurskainen, 1996; Perry-O'Keefe *et al.*, 1996).

High-affinity binding provides clear advantages for molecular recognition and the development of new applications for PNAs. For example, 11-13 nucleotide PNAs inhibit the activity of telomerase, a ribonucleo-protein that extends telomere ends using an

15 essential RNA template, while the analogous DNA oligomers do not (Norton *et al.*, 1996).

Neutral PNAs are more hydrophobic than analogous DNA oligomers, and this can lead to difficulty solubilizing them at neutral pH, especially if the PNAs have a high purine content or if they have the potential to form secondary structures. Their solubility can be enhanced by attaching one or more positive charges to the PNA termini

20 (Nielsen *et al.*, 1991).

Findings by Allfrey and colleagues suggest that strand invasion will occur spontaneously at sequences within chromosomal DNA (Boffa *et al.*, 1995; Boffa *et al.*, 1996). These studies targeted PNAs to triplet repeats of the nucleotides CAG and used this recognition to purify transcriptionally active DNA (Boffa *et al.*, 1995) and to inhibit

25 transcription (Boffa *et al.*, 1996). This result suggests that if PNAs can be delivered within cells then they will have the potential to be general sequence-specific regulators of gene expression. Studies and reviews concerning the use of PNAs as antisense and anti-gene agents include Nielsen *et al.* (1993b), Hanvey *et al.* (1992), and Good and Nielsen (1997).

Koppelhus *et al.* (1997) have used PNAs to inhibit HIV-1 inverse transcription, showing that PNAs may be used for antiviral therapies.

Methods of characterizing the antisense binding properties of PNAs are discussed in Rose (1993) and Jensen *et al.* (1997). Rose uses capillary gel electrophoresis to determine binding of PNAs to their complementary oligonucleotide, measuring the relative binding kinetics and stoichiometry. Similar types of measurements were made by Jensen *et al.* using BIAcore™ technology.

Other applications of PNAs include use in DNA strand invasion (Nielsen *et al.*, 1991), antisense inhibition (Hanvey *et al.*, 1992), mutational analysis (Orum *et al.*, 1993), enhancers of transcription (Mollegaard *et al.*, 1994), nucleic acid purification (Orum *et al.*, 1995), isolation of transcriptionally active genes (Boffa *et al.*, 1995), blocking of transcription factor binding (Vickers *et al.*, 1995), genome cleavage (Veselkov *et al.*, 1996), biosensors (Wang *et al.*, 1996), *in situ* hybridization (Thisted *et al.*, 1996), and in an alternative to Southern blotting (Perry-O'Keefe, 1996).

15 POLYPEPTIDE COMPOSITIONS

The present invention, in other aspects, provides polypeptide compositions. Generally, a polypeptide of the invention will be an isolated polypeptide (or an epitope, variant, or active fragment thereof) derived from a mammalian species. Preferably, the polypeptide is encoded by a polynucleotide sequence disclosed herein or a sequence which hybridizes under moderately stringent conditions to a polynucleotide sequence disclosed herein. Alternatively, the polypeptide may be defined as a polypeptide which comprises a contiguous amino acid sequence from an amino acid sequence disclosed herein, or which polypeptide comprises an entire amino acid sequence disclosed herein.

In the present invention, a polypeptide composition is also understood to comprise one or more polypeptides that are immunologically reactive with antibodies generated against a polypeptide of the invention, particularly a polypeptide having the amino acid sequence disclosed in SEQ ID NO: 786, 787, 791, 793, 795, 797-799, 806, 809,

1670-1675, or to active fragments, or to variants or biological functional equivalents thereof.

Likewise, a polypeptide composition of the present invention is understood to comprise one or more polypeptides that are capable of eliciting antibodies that are immunologically reactive with one or more polypeptides encoded by one or more contiguous nucleic acid sequences contained in SEQ ID NO: 1, 11-13, 15, 20, 23-27, 29, 30, 33, 34, 39, 41, 43-46, 51, 52, 57, 58, 60, 62, 65-67, 69-71, 74, 76, 79, 80, 84, 86, 89-92, 95, 97, 98, 101, 110, 111, 113-119, 121-128, 130-134, 136, 138, 139, 141, 143, 146-151, 153, 154, 157-160, 162-164, 167-178, 180, 181, 183, 186-190, 192, 193, 195-220, 224, 226-231, 234, 236, 237, 240, 241, 244-246, 248, 254, 255, 261, 262, 266, 270, 275, 280, 282, 283, 288, 289, 290, 292, 295, 301, 303, 304, 309, 311, 341-782, 784, 785, 790, 792, 794, 796, 800-804, 807, 808, 810-826, 828-1664, 1669 and 1676 or to active fragments, or to variants thereof, or to one or more nucleic acid sequences which hybridize to one or more of these sequences under conditions of moderate to high stringency. Particularly illustrative polypeptides include the amino acid sequences disclosed in SEQ ID NO: 786, 787, 791, 793, 795, 797-799, 806, 809, 827 and 1670-1675..

As used herein, an active fragment of a polypeptide includes a whole or a portion of a polypeptide which is modified by conventional techniques, *e.g.*, mutagenesis, or by addition, deletion, or substitution, but which active fragment exhibits substantially the same structure function, antigenicity, etc., as a polypeptide as described herein.

In certain illustrative embodiments, the polypeptides of the invention will comprise at least an immunogenic portion of a lung tumor protein or a variant thereof, as described herein. As noted above, a "lung tumor protein" is a protein that is expressed by lung tumor cells. Proteins that are lung tumor proteins also react detectably within an immunoassay (such as an ELISA) with antisera from a patient with lung cancer. Polypeptides as described herein may be of any length. Additional sequences derived from the native protein and/or heterologous sequences may be present, and such sequences may (but need not) possess further immunogenic or antigenic properties.

An "immunogenic portion," as used herein is a portion of a protein that is recognized (*i.e.*, specifically bound) by a B-cell and/or T-cell surface antigen receptor. Such immunogenic portions generally comprise at least 5 amino acid residues, more preferably at least 10, and still more preferably at least 20 amino acid residues of a lung tumor protein or a variant thereof. Certain preferred immunogenic portions include peptides in which an N-terminal leader sequence and/or transmembrane domain have been deleted. Other preferred immunogenic portions may contain a small N- and/or C-terminal deletion (*e.g.*, 1-30 amino acids, preferably 5-15 amino acids), relative to the mature protein.

Immunogenic portions may generally be identified using well known techniques, such as those summarized in Paul, *Fundamental Immunology*, 3rd ed., 243-247 (Raven Press, 1993) and references cited therein. Such techniques include screening polypeptides for the ability to react with antigen-specific antibodies, antisera and/or T-cell lines or clones. As used herein, antisera and antibodies are "antigen-specific" if they specifically bind to an antigen (*i.e.*, they react with the protein in an ELISA or other immunoassay, and do not react detectably with unrelated proteins). Such antisera and antibodies may be prepared as described herein, and using well known techniques. An immunogenic portion of a native lung tumor protein is a portion that reacts with such antisera and/or T-cells at a level that is not substantially less than the reactivity of the full length polypeptide (*e.g.*, in an ELISA and/or T-cell reactivity assay). Such immunogenic portions may react within such assays at a level that is similar to or greater than the reactivity of the full length polypeptide. Such screens may generally be performed using methods well known to those of ordinary skill in the art, such as those described in Harlow and Lane, *Antibodies: A Laboratory Manual*, Cold Spring Harbor Laboratory, 1988. For example, a polypeptide may be immobilized on a solid support and contacted with patient sera to allow binding of antibodies within the sera to the immobilized polypeptide. Unbound sera may then be removed and bound antibodies detected using, for example, ¹²⁵I-labeled Protein A.

As noted above, a composition may comprise a variant of a native lung tumor protein. A polypeptide "variant," as used herein, is a polypeptide that differs from a native lung tumor protein in one or more substitutions, deletions, additions and/or insertions, such that the immunogenicity of the polypeptide is not substantially diminished.

5 In other words, the ability of a variant to react with antigen-specific antisera may be enhanced or unchanged, relative to the native protein, or may be diminished by less than 50%, and preferably less than 20%, relative to the native protein. Such variants may generally be identified by modifying one of the above polypeptide sequences and evaluating the reactivity of the modified polypeptide with antigen-specific antibodies or
10 antisera as described herein. Preferred variants include those in which one or more portions, such as an N-terminal leader sequence or transmembrane domain, have been removed. Other preferred variants include variants in which a small portion (*e.g.*, 1-30 amino acids, preferably 5-15 amino acids) has been removed from the N- and/or C-terminal of the mature protein.

15 Polypeptide variants encompassed by the present invention include those exhibiting at least about 70%, 75%, 80%, 85%, 90%, 91%, 92%, 93%, 94%, 95%, 96%, 97%, 98%, or 99% or more identity (determined as described above) to the polypeptides disclosed herein.

Preferably, a variant contains conservative substitutions. A "conservative
20 substitution" is one in which an amino acid is substituted for another amino acid that has similar properties, such that one skilled in the art of peptide chemistry would expect the secondary structure and hydrophobic nature of the polypeptide to be substantially unchanged. Amino acid substitutions may generally be made on the basis of similarity in polarity, charge, solubility, hydrophobicity, hydrophilicity and/or the amphipathic nature of
25 the residues. For example, negatively charged amino acids include aspartic acid and glutamic acid; positively charged amino acids include lysine and arginine; and amino acids with uncharged polar head groups having similar hydrophilicity values include leucine, isoleucine and valine; glycine and alanine; asparagine and glutamine; and serine, threonine, phenylalanine and tyrosine. Other groups of amino acids that may represent conservative

changes include: (1) ala, pro, gly, glu, asp, gln, asn, ser, thr; (2) cys, ser, tyr, thr; (3) val, ile, leu, met, ala, phe; (4) lys, arg, his; and (5) phe, tyr, trp, his. A variant may also, or alternatively, contain nonconservative changes. In a preferred embodiment, variant polypeptides differ from a native sequence by substitution, deletion or addition of five
 5 amino acids or fewer. Variants may also (or alternatively) be modified by, for example, the deletion or addition of amino acids that have minimal influence on the immunogenicity, secondary structure and hydrophobic nature of the polypeptide.

As noted above, polypeptides may comprise a signal (or leader) sequence at the N-terminal end of the protein, which co-translationally or post-translationally directs
 10 transfer of the protein. The polypeptide may also be conjugated to a linker or other sequence for ease of synthesis, purification or identification of the polypeptide (*e.g.*, poly-His), or to enhance binding of the polypeptide to a solid support. For example, a polypeptide may be conjugated to an immunoglobulin Fc region.

Polypeptides may be prepared using any of a variety of well known
 15 techniques. Recombinant polypeptides encoded by DNA sequences as described above may be readily prepared from the DNA sequences using any of a variety of expression vectors known to those of ordinary skill in the art. Expression may be achieved in any appropriate host cell that has been transformed or transfected with an expression vector containing a DNA molecule that encodes a recombinant polypeptide. Suitable host cells
 20 include prokaryotes, yeast, and higher eukaryotic cells, such as mammalian cells and plant cells. Preferably, the host cells employed are *E. coli*, yeast or a mammalian cell line such as COS or CHO. Supernatants from suitable host/vector systems which secrete recombinant protein or polypeptide into culture media may be first concentrated using a commercially available filter. Following concentration, the concentrate may be applied to
 25 a suitable purification matrix such as an affinity matrix or an ion exchange resin. Finally, one or more reverse phase HPLC steps can be employed to further purify a recombinant polypeptide.

Portions and other variants having less than about 100 amino acids, and generally less than about 50 amino acids, may also be generated by synthetic means, using

techniques well known to those of ordinary skill in the art. For example, such polypeptides may be synthesized using any of the commercially available solid-phase techniques, such as the Merrifield solid-phase synthesis method, where amino acids are sequentially added to a growing amino acid chain. *See* Merrifield, *J. Am. Chem. Soc.* 85:2149-2146, 1963.

- 5 Equipment for automated synthesis of polypeptides is commercially available from suppliers such as Perkin Elmer/Applied BioSystems Division (Foster City, CA), and may be operated according to the manufacturer's instructions.

Within certain specific embodiments, a polypeptide may be a fusion protein that comprises multiple polypeptides as described herein, or that comprises at least one
 10 polypeptide as described herein and an unrelated sequence, such as a known tumor protein. A fusion partner may, for example, assist in providing T helper epitopes (an immunological fusion partner), preferably T helper epitopes recognized by humans, or may assist in expressing the protein (an expression enhancer) at higher yields than the native recombinant protein. Certain preferred fusion partners are both immunological and
 15 expression enhancing fusion partners. Other fusion partners may be selected so as to increase the solubility of the protein or to enable the protein to be targeted to desired intracellular compartments. Still further fusion partners include affinity tags, which facilitate purification of the protein.

Fusion proteins may generally be prepared using standard techniques,
 20 including chemical conjugation. Preferably, a fusion protein is expressed as a recombinant protein, allowing the production of increased levels, relative to a non-fused protein, in an expression system. Briefly, DNA sequences encoding the polypeptide components may be assembled separately, and ligated into an appropriate expression vector. The 3' end of the DNA sequence encoding one polypeptide component is ligated, with or without a peptide
 25 linker, to the 5' end of a DNA sequence encoding the second polypeptide component so that the reading frames of the sequences are in phase. This permits translation into a single fusion protein that retains the biological activity of both component polypeptides.

A peptide linker sequence may be employed to separate the first and second polypeptide components by a distance sufficient to ensure that each polypeptide folds into

its secondary and tertiary structures. Such a peptide linker sequence is incorporated into the fusion protein using standard techniques well known in the art. Suitable peptide linker sequences may be chosen based on the following factors: (1) their ability to adopt a flexible extended conformation; (2) their inability to adopt a secondary structure that could interact with functional epitopes on the first and second polypeptides; and (3) the lack of hydrophobic or charged residues that might react with the polypeptide functional epitopes. Preferred peptide linker sequences contain Gly, Asn and Ser residues. Other near neutral amino acids, such as Thr and Ala may also be used in the linker sequence. Amino acid sequences which may be usefully employed as linkers include those disclosed in Maratea *et al.*, *Gene* 40:39-46, 1985; Murphy *et al.*, *Proc. Natl. Acad. Sci. USA* 83:8258-8262, 1986; U.S. Patent No. 4,935,233 and U.S. Patent No. 4,751,180. The linker sequence may generally be from 1 to about 50 amino acids in length. Linker sequences are not required when the first and second polypeptides have non-essential N-terminal amino acid regions that can be used to separate the functional domains and prevent steric interference.

The ligated DNA sequences are operably linked to suitable transcriptional or translational regulatory elements. The regulatory elements responsible for expression of DNA are located only 5' to the DNA sequence encoding the first polypeptides. Similarly, stop codons required to end translation and transcription termination signals are only present 3' to the DNA sequence encoding the second polypeptide.

Fusion proteins are also provided. Such proteins comprise a polypeptide as described herein together with an unrelated immunogenic protein. Preferably the immunogenic protein is capable of eliciting a recall response. Examples of such proteins include tetanus, tuberculosis and hepatitis proteins (*see*, for example, Stoute *et al.* *New Engl. J. Med.*, 336:86-91, 1997).

Within preferred embodiments, an immunological fusion partner is derived from protein D, a surface protein of the gram-negative bacterium *Haemophilus influenza* B (WO 91/18926). Preferably, a protein D derivative comprises approximately the first third of the protein (*e.g.*, the first N-terminal 100-110 amino acids), and a protein D derivative may be lipidated. Within certain preferred embodiments, the first 109 residues of a

Lipoprotein D fusion partner is included on the N-terminus to provide the polypeptide with additional exogenous T-cell epitopes and to increase the expression level in *E. coli* (thus functioning as an expression enhancer). The lipid tail ensures optimal presentation of the antigen to antigen presenting cells. Other fusion partners include the non-structural protein from influenzae virus, NS1 (hemagglutinin). Typically, the N-terminal 81 amino acids are used, although different fragments that include T-helper epitopes may be used.

In another embodiment, the immunological fusion partner is the protein known as LYTA, or a portion thereof (preferably a C-terminal portion). LYTA is derived from *Streptococcus pneumoniae*, which synthesizes an N-acetyl-L-alanine amidase known as amidase LYTA (encoded by the *LytA* gene; *Gene* 43:265-292, 1986). LYTA is an autolysin that specifically degrades certain bonds in the peptidoglycan backbone. The C-terminal domain of the LYTA protein is responsible for the affinity to the choline or to some choline analogues such as DEAE. This property has been exploited for the development of *E. coli* C-LYTA expressing plasmids useful for expression of fusion proteins. Purification of hybrid proteins containing the C-LYTA fragment at the amino terminus has been described (*see Biotechnology* 10:795-798, 1992). Within a preferred embodiment, a repeat portion of LYTA may be incorporated into a fusion protein. A repeat portion is found in the C-terminal region starting at residue 178. A particularly preferred repeat portion incorporates residues 188-305.

In general, polypeptides (including fusion proteins) and polynucleotides as described herein are isolated. An "isolated" polypeptide or polynucleotide is one that is removed from its original environment. For example, a naturally-occurring protein is isolated if it is separated from some or all of the coexisting materials in the natural system. Preferably, such polypeptides are at least about 90% pure, more preferably at least about 95% pure and most preferably at least about 99% pure. A polynucleotide is considered to be isolated if, for example, it is cloned into a vector that is not a part of the natural environment.

BINDING AGENTS

The present invention further provides agents, such as antibodies and antigen-binding fragments thereof, that specifically bind to a lung tumor protein. As used herein, an antibody, or antigen-binding fragment thereof, is said to "specifically bind" to a lung tumor protein if it reacts at a detectable level (within, for example, an ELISA) with a lung tumor protein, and does not react detectably with unrelated proteins under similar conditions. As used herein, "binding" refers to a noncovalent association between two separate molecules such that a complex is formed. The ability to bind may be evaluated by, for example, determining a binding constant for the formation of the complex. The binding constant is the value obtained when the concentration of the complex is divided by the product of the component concentrations. In general, two compounds are said to "bind," in the context of the present invention, when the binding constant for complex formation exceeds about 10^3 L/mol. The binding constant may be determined using methods well known in the art.

Binding agents may be further capable of differentiating between patients with and without a cancer, such as lung cancer, using the representative assays provided herein. In other words, antibodies or other binding agents that bind to a lung tumor protein will generate a signal indicating the presence of a cancer in at least about 20% of patients with the disease, and will generate a negative signal indicating the absence of the disease in at least about 90% of individuals without the cancer. To determine whether a binding agent satisfies this requirement, biological samples (*e.g.*, blood, sera, sputum, urine and/or tumor biopsies) from patients with and without a cancer (as determined using standard clinical tests) may be assayed as described herein for the presence of polypeptides that bind to the binding agent. It will be apparent that a statistically significant number of samples with and without the disease should be assayed. Each binding agent should satisfy the above criteria; however, those of ordinary skill in the art will recognize that binding agents may be used in combination to improve sensitivity.

Any agent that satisfies the above requirements may be a binding agent. For example, a binding agent may be a ribosome, with or without a peptide component, an

RNA molecule or a polypeptide. In a preferred embodiment, a binding agent is an antibody or an antigen-binding fragment thereof. Antibodies may be prepared by any of a variety of techniques known to those of ordinary skill in the art. *See, e.g.,* Harlow and Lane, *Antibodies: A Laboratory Manual*, Cold Spring Harbor Laboratory, 1988. In
5 general, antibodies can be produced by cell culture techniques, including the generation of monoclonal antibodies as described herein, or via transfection of antibody genes into suitable bacterial or mammalian cell hosts, in order to allow for the production of recombinant antibodies. In one technique, an immunogen comprising the polypeptide is initially injected into any of a wide variety of mammals (*e.g.,* mice, rats, rabbits, sheep or
10 goats). In this step, the polypeptides of this invention may serve as the immunogen without modification. Alternatively, particularly for relatively short polypeptides, a superior immune response may be elicited if the polypeptide is joined to a carrier protein, such as bovine serum albumin or keyhole limpet hemocyanin. The immunogen is injected into the animal host, preferably according to a predetermined schedule incorporating one or more
15 booster immunizations, and the animals are bled periodically. Polyclonal antibodies specific for the polypeptide may then be purified from such antisera by, for example, affinity chromatography using the polypeptide coupled to a suitable solid support.

Monoclonal antibodies specific for an antigenic polypeptide of interest may be prepared, for example, using the technique of Kohler and Milstein, *Eur. J. Immunol.*
20 6:511-519, 1976, and improvements thereto. Briefly, these methods involve the preparation of immortal cell lines capable of producing antibodies having the desired specificity (*i.e.,* reactivity with the polypeptide of interest). Such cell lines may be produced, for example, from spleen cells obtained from an animal immunized as described above. The spleen cells are then immortalized by, for example, fusion with a myeloma cell
25 fusion partner, preferably one that is syngeneic with the immunized animal. A variety of fusion techniques may be employed. For example, the spleen cells and myeloma cells may be combined with a nonionic detergent for a few minutes and then plated at low density on a selective medium that supports the growth of hybrid cells, but not myeloma cells. A preferred selection technique uses HAT (hypoxanthine, aminopterin, thymidine) selection.

After a sufficient time, usually about 1 to 2 weeks, colonies of hybrids are observed. Single colonies are selected and their culture supernatants tested for binding activity against the polypeptide. Hybridomas having high reactivity and specificity are preferred.

Monoclonal antibodies may be isolated from the supernatants of growing hybridoma colonies. In addition, various techniques may be employed to enhance the yield, such as injection of the hybridoma cell line into the peritoneal cavity of a suitable vertebrate host, such as a mouse. Monoclonal antibodies may then be harvested from the ascites fluid or the blood. Contaminants may be removed from the antibodies by conventional techniques, such as chromatography, gel filtration, precipitation, and extraction. The polypeptides of this invention may be used in the purification process in, for example, an affinity chromatography step.

Within certain embodiments, the use of antigen-binding fragments of antibodies may be preferred. Such fragments include Fab fragments, which may be prepared using standard techniques. Briefly, immunoglobulins may be purified from rabbit serum by affinity chromatography on Protein A bead columns (Harlow and Lane, *Antibodies: A Laboratory Manual*, Cold Spring Harbor Laboratory, 1988) and digested by papain to yield Fab and Fc fragments. The Fab and Fc fragments may be separated by affinity chromatography on protein A bead columns.

Monoclonal antibodies of the present invention may be coupled to one or more therapeutic agents. Suitable agents in this regard include radionuclides, differentiation inducers, drugs, toxins, and derivatives thereof. Preferred radionuclides include ^{90}Y , ^{123}I , ^{125}I , ^{131}I , ^{186}Re , ^{188}Re , ^{211}At , and ^{212}Bi . Preferred drugs include methotrexate, and pyrimidine and purine analogs. Preferred differentiation inducers include phorbol esters and butyric acid. Preferred toxins include ricin, abrin, diphtheria toxin, cholera toxin, gelonin, *Pseudomonas* exotoxin, *Shigella* toxin, and pokeweed antiviral protein.

A therapeutic agent may be coupled (*e.g.*, covalently bonded) to a suitable monoclonal antibody either directly or indirectly (*e.g.*, via a linker group). A direct reaction between an agent and an antibody is possible when each possesses a substituent

capable of reacting with the other. For example, a nucleophilic group, such as an amino or sulfhydryl group, on one may be capable of reacting with a carbonyl-containing group, such as an anhydride or an acid halide, or with an alkyl group containing a good leaving group (*e.g.*, a halide) on the other.

5 Alternatively, it may be desirable to couple a therapeutic agent and an antibody via a linker group. A linker group can function as a spacer to distance an antibody from an agent in order to avoid interference with binding capabilities. A linker group can also serve to increase the chemical reactivity of a substituent on an agent or an antibody, and thus increase the coupling efficiency. An increase in chemical reactivity
10 may also facilitate the use of agents, or functional groups on agents, which otherwise would not be possible.

 It will be evident to those skilled in the art that a variety of bifunctional or polyfunctional reagents, both homo- and hetero-functional (such as those described in the catalog of the Pierce Chemical Co., Rockford, IL), may be employed as the linker group.
15 Coupling may be effected, for example, through amino groups, carboxyl groups, sulfhydryl groups or oxidized carbohydrate residues. There are numerous references describing such methodology, *e.g.*, U.S. Patent No. 4,671,958, to Rodwell *et al.*

 Where a therapeutic agent is more potent when free from the antibody portion of the immunoconjugates of the present invention, it may be desirable to use a
20 linker group which is cleavable during or upon internalization into a cell. A number of different cleavable linker groups have been described. The mechanisms for the intracellular release of an agent from these linker groups include cleavage by reduction of a disulfide bond (*e.g.*, U.S. Patent No. 4,489,710, to Spitler), by irradiation of a photolabile bond (*e.g.*, U.S. Patent No. 4,625,014, to Senter *et al.*), by hydrolysis of derivatized amino
25 acid side chains (*e.g.*, U.S. Patent No. 4,638,045, to Kohn *et al.*), by serum complement-mediated hydrolysis (*e.g.*, U.S. Patent No. 4,671,958, to Rodwell *et al.*), and acid-catalyzed hydrolysis (*e.g.*, U.S. Patent No. 4,569,789, to Blattler *et al.*).

 It may be desirable to couple more than one agent to an antibody. In one embodiment, multiple molecules of an agent are coupled to one antibody molecule. In

another embodiment, more than one type of agent may be coupled to one antibody. Regardless of the particular embodiment, immunoconjugates with more than one agent may be prepared in a variety of ways. For example, more than one agent may be coupled directly to an antibody molecule, or linkers that provide multiple sites for attachment can
 5 be used. Alternatively, a carrier can be used.

A carrier may bear the agents in a variety of ways, including covalent bonding either directly or via a linker group. Suitable carriers include proteins such as albumins (*e.g.*, U.S. Patent No. 4,507,234, to Kato *et al.*), peptides and polysaccharides such as aminodextran (*e.g.*, U.S. Patent No. 4,699,784, to Shih *et al.*). A carrier may also
 10 bear an agent by noncovalent bonding or by encapsulation, such as within a liposome vesicle (*e.g.*, U.S. Patent Nos. 4,429,008 and 4,873,088). Carriers specific for radionuclide agents include radiohalogenated small molecules and chelating compounds. For example, U.S. Patent No. 4,735,792 discloses representative radiohalogenated small molecules and their synthesis. A radionuclide chelate may be formed from chelating compounds that
 15 include those containing nitrogen and sulfur atoms as the donor atoms for binding the metal, or metal oxide, radionuclide. For example, U.S. Patent No. 4,673,562, to Davison *et al.* discloses representative chelating compounds and their synthesis.

A variety of routes of administration for the antibodies and immunoconjugates may be used. Typically, administration will be intravenous,
 20 intramuscular, subcutaneous or in the bed of a resected tumor. It will be evident that the precise dose of the antibody/immunoconjugate will vary depending upon the antibody used, the antigen density on the tumor, and the rate of clearance of the antibody.

T CELLS

Immunotherapeutic compositions may also, or alternatively, comprise T
 25 cells specific for a lung tumor protein. Such cells may generally be prepared *in vitro* or *ex vivo*, using standard procedures. For example, T cells may be isolated from bone marrow, peripheral blood, or a fraction of bone marrow or peripheral blood of a patient, using a commercially available cell separation system, such as the Isolex™ System, available from

Nexell Therapeutics, Inc. (Irvine, CA; see also U.S. Patent No. 5,240,856; U.S. Patent No. 5,215,926; WO 89/06280; WO 91/16116 and WO 92/07243). Alternatively, T cells may be derived from related or unrelated humans, non-human mammals, cell lines or cultures.

5 T cells may be stimulated with a lung tumor polypeptide, polynucleotide encoding a lung tumor polypeptide and/or an antigen presenting cell (APC) that expresses such a polypeptide. Such stimulation is performed under conditions and for a time sufficient to permit the generation of T cells that are specific for the polypeptide. Preferably, a lung tumor polypeptide or polynucleotide is present within a delivery vehicle, such as a microsphere, to facilitate the generation of specific T cells.

10 T cells are considered to be specific for a lung tumor polypeptide if the T cells specifically proliferate, secrete cytokines or kill target cells coated with the polypeptide or expressing a gene encoding the polypeptide. T cell specificity may be evaluated using any of a variety of standard techniques. For example, within a chromium release assay or proliferation assay, a stimulation index of more than two fold increase in
15 lysis and/or proliferation, compared to negative controls, indicates T cell specificity. Such assays may be performed, for example, as described in Chen *et al.*, *Cancer Res.* 54:1065-1070, 1994. Alternatively, detection of the proliferation of T cells may be accomplished by a variety of known techniques. For example, T cell proliferation can be detected by measuring an increased rate of DNA synthesis (*e.g.*, by pulse-labeling cultures of T cells
20 with tritiated thymidine and measuring the amount of tritiated thymidine incorporated into DNA). Contact with a lung tumor polypeptide (100 ng/ml - 100 µg/ml, preferably 200 ng/ml - 25 µg/ml) for 3 - 7 days should result in at least a two fold increase in proliferation of the T cells. Contact as described above for 2-3 hours should result in activation of the T cells, as measured using standard cytokine assays in which a two fold increase in the level
25 of cytokine release (*e.g.*, TNF or IFN-γ) is indicative of T cell activation (*see* Coligan *et al.*, *Current Protocols in Immunology*, vol. 1, Wiley Interscience (Greene 1998)). T cells that have been activated in response to a lung tumor polypeptide, polynucleotide or polypeptide-expressing APC may be CD4⁺ and/or CD8⁺. Lung tumor protein-specific T cells may be expanded using standard techniques. Within preferred embodiments, the T

cells are derived from a patient, a related donor or an unrelated donor, and are administered to the patient following stimulation and expansion.

For therapeutic purposes, CD4⁺ or CD8⁺ T cells that proliferate in response to a lung tumor polypeptide, polynucleotide or APC can be expanded in number either *in vitro* or *in vivo*. Proliferation of such T cells *in vitro* may be accomplished in a variety of ways. For example, the T cells can be re-exposed to a lung tumor polypeptide, or a short peptide corresponding to an immunogenic portion of such a polypeptide, with or without the addition of T cell growth factors, such as interleukin-2, and/or stimulator cells that synthesize a lung tumor polypeptide. Alternatively, one or more T cells that proliferate in the presence of a lung tumor protein can be expanded in number by cloning. Methods for cloning cells are well known in the art, and include limiting dilution.

PHARMACEUTICAL COMPOSITIONS

In additional embodiments, the present invention concerns formulation of one or more of the polynucleotide, polypeptide, T-cell and/or antibody compositions disclosed herein in pharmaceutically-acceptable solutions for administration to a cell or an animal, either alone, or in combination with one or more other modalities of therapy.

It will also be understood that, if desired, the nucleic acid segment, RNA, DNA or PNA compositions that express a polypeptide as disclosed herein may be administered in combination with other agents as well, such as, *e.g.*, other proteins or polypeptides or various pharmaceutically-active agents. In fact, there is virtually no limit to other components that may also be included, given that the additional agents do not cause a significant adverse effect upon contact with the target cells or host tissues. The compositions may thus be delivered along with various other agents as required in the particular instance. Such compositions may be purified from host cells or other biological sources, or alternatively may be chemically synthesized as described herein. Likewise, such compositions may further comprise substituted or derivatized RNA or DNA compositions.

Formulation of pharmaceutically-acceptable excipients and carrier solutions is well-known to those of skill in the art, as is the development of suitable dosing and treatment regimens for using the particular compositions described herein in a variety of treatment regimens, including *e.g.*, oral, parenteral, intravenous, intranasal, and intramuscular administration and formulation.

1. ORAL DELIVERY

In certain applications, the pharmaceutical compositions disclosed herein may be delivered *via* oral administration to an animal. As such, these compositions may be formulated with an inert diluent or with an assimilable edible carrier, or they may be enclosed in hard- or soft-shell gelatin capsule, or they may be compressed into tablets, or they may be incorporated directly with the food of the diet.

The active compounds may even be incorporated with excipients and used in the form of ingestible tablets, buccal tables, troches, capsules, elixirs, suspensions, syrups, wafers, and the like (Mathiowitz *et al.*, 1997; Hwang *et al.*, 1998; U. S. Patent 5,641,515; U. S. Patent 5,580,579 and U. S. Patent 5,792,451, each specifically incorporated herein by reference in its entirety). The tablets, troches, pills, capsules and the like may also contain the following: a binder, as gum tragacanth, acacia, cornstarch, or gelatin; excipients, such as dicalcium phosphate; a disintegrating agent, such as corn starch, potato starch, alginic acid and the like; a lubricant, such as magnesium stearate; and a sweetening agent, such as sucrose, lactose or saccharin may be added or a flavoring agent, such as peppermint, oil of wintergreen, or cherry flavoring. When the dosage unit form is a capsule, it may contain, in addition to materials of the above type, a liquid carrier. Various other materials may be present as coatings or to otherwise modify the physical form of the dosage unit. For instance, tablets, pills, or capsules may be coated with shellac, sugar, or both. A syrup or elixir may contain the active compound sucrose as a sweetening agent methyl and propylparabens as preservatives, a dye and flavoring, such as cherry or orange flavor. Of course, any material used in preparing any dosage unit form should be pharmaceutically pure and substantially non-toxic in the amounts employed. In addition,

the active compounds may be incorporated into sustained-release preparation and formulations.

Typically, these formulations may contain at least about 0.1% of the active compound or more, although the percentage of the active ingredient(s) may, of course, be varied and may conveniently be between about 1 or 2% and about 60% or 70% or more of the weight or volume of the total formulation. Naturally, the amount of active compound(s) in each therapeutically useful composition may be prepared in such a way that a suitable dosage will be obtained in any given unit dose of the compound. Factors such as solubility, bioavailability, biological half-life, route of administration, product shelf life, as well as other pharmacological considerations will be contemplated by one skilled in the art of preparing such pharmaceutical formulations, and as such, a variety of dosages and treatment regimens may be desirable.

For oral administration the compositions of the present invention may alternatively be incorporated with one or more excipients in the form of a mouthwash, dentifrice, buccal tablet, oral spray, or sublingual orally-administered formulation. For example, a mouthwash may be prepared incorporating the active ingredient in the required amount in an appropriate solvent, such as a sodium borate solution (Dobell's Solution). Alternatively, the active ingredient may be incorporated into an oral solution such as one containing sodium borate, glycerin and potassium bicarbonate, or dispersed in a dentifrice, or added in a therapeutically-effective amount to a composition that may include water, binders, abrasives, flavoring agents, foaming agents, and humectants. Alternatively the compositions may be fashioned into a tablet or solution form that may be placed under the tongue or otherwise dissolved in the mouth.

2. INJECTABLE DELIVERY

In certain circumstances it will be desirable to deliver the pharmaceutical compositions disclosed herein parenterally, intravenously, intramuscularly, or even intraperitoneally as described in U. S. Patent 5,543,158; U. S. Patent 5,641,515 and U. S. Patent 5,399,363 (each specifically incorporated herein by reference in its entirety).

Solutions of the active compounds as free base or pharmacologically acceptable salts may be prepared in water suitably mixed with a surfactant, such as hydroxypropylcellulose. Dispersions may also be prepared in glycerol, liquid polyethylene glycols, and mixtures thereof and in oils. Under ordinary conditions of storage and use, these preparations
5 contain a preservative to prevent the growth of microorganisms.

The pharmaceutical forms suitable for injectable use include sterile aqueous solutions or dispersions and sterile powders for the extemporaneous preparation of sterile injectable solutions or dispersions (U. S. Patent 5,466,468, specifically incorporated herein by reference in its entirety). In all cases the form must be sterile and must be fluid to the
10 extent that easy syringability exists. It must be stable under the conditions of manufacture and storage and must be preserved against the contaminating action of microorganisms, such as bacteria and fungi. The carrier can be a solvent or dispersion medium containing, for example, water, ethanol, polyol (*e.g.*, glycerol, propylene glycol, and liquid polyethylene glycol, and the like), suitable mixtures thereof, and/or vegetable oils. Proper
15 fluidity may be maintained, for example, by the use of a coating, such as lecithin, by the maintenance of the required particle size in the case of dispersion and by the use of surfactants. The prevention of the action of microorganisms can be facilitated by various antibacterial and antifungal agents, for example, parabens, chlorobutanol, phenol, sorbic acid, thimerosal, and the like. In many cases, it will be preferable to include isotonic
20 agents, for example, sugars or sodium chloride. Prolonged absorption of the injectable compositions can be brought about by the use in the compositions of agents delaying absorption, for example, aluminum monostearate and gelatin.

For parenteral administration in an aqueous solution, for example, the solution should be suitably buffered if necessary and the liquid diluent first rendered
25 isotonic with sufficient saline or glucose. These particular aqueous solutions are especially suitable for intravenous, intramuscular, subcutaneous and intraperitoneal administration. In this connection, a sterile aqueous medium that can be employed will be known to those of skill in the art in light of the present disclosure. For example, one dosage may be dissolved in 1 ml of isotonic NaCl solution and either added to 1000 ml of hypodermoclysis fluid or

injected at the proposed site of infusion, (see for example, "Remington's Pharmaceutical Sciences" 15th Edition, pages 1035-1038 and 1570-1580). Some variation in dosage will necessarily occur depending on the condition of the subject being treated. The person responsible for administration will, in any event, determine the appropriate dose for the individual subject. Moreover, for human administration, preparations should meet sterility, pyrogenicity, and the general safety and purity standards as required by FDA Office of Biologics standards.

Sterile injectable solutions are prepared by incorporating the active compounds in the required amount in the appropriate solvent with various of the other ingredients enumerated above, as required, followed by filtered sterilization. Generally, dispersions are prepared by incorporating the various sterilized active ingredients into a sterile vehicle which contains the basic dispersion medium and the required other ingredients from those enumerated above. In the case of sterile powders for the preparation of sterile injectable solutions, the preferred methods of preparation are vacuum-drying and freeze-drying techniques which yield a powder of the active ingredient plus any additional desired ingredient from a previously sterile-filtered solution thereof.

The compositions disclosed herein may be formulated in a neutral or salt form. Pharmaceutically-acceptable salts, include the acid addition salts (formed with the free amino groups of the protein) and which are formed with inorganic acids such as, for example, hydrochloric or phosphoric acids, or such organic acids as acetic, oxalic, tartaric, mandelic, and the like. Salts formed with the free carboxyl groups can also be derived from inorganic bases such as, for example, sodium, potassium, ammonium, calcium, or ferric hydroxides, and such organic bases as isopropylamine, trimethylamine, histidine, procaine and the like. Upon formulation, solutions will be administered in a manner compatible with the dosage formulation and in such amount as is therapeutically effective. The formulations are easily administered in a variety of dosage forms such as injectable solutions, drug-release capsules, and the like.

As used herein, "carrier" includes any and all solvents, dispersion media, vehicles, coatings, diluents, antibacterial and antifungal agents, isotonic and absorption

delaying agents, buffers, carrier solutions, suspensions, colloids, and the like. The use of such media and agents for pharmaceutical active substances is well known in the art. Except insofar as any conventional media or agent is incompatible with the active ingredient, its use in the therapeutic compositions is contemplated. Supplementary active
 5 ingredients can also be incorporated into the compositions.

The phrase "pharmaceutically-acceptable" refers to molecular entities and compositions that do not produce an allergic or similar untoward reaction when administered to a human. The preparation of an aqueous composition that contains a protein as an active ingredient is well understood in the art. Typically, such compositions
 10 are prepared as injectables, either as liquid solutions or suspensions; solid forms suitable for solution in, or suspension in, liquid prior to injection can also be prepared. The preparation can also be emulsified.

3. NASAL DELIVERY

In certain embodiments, the pharmaceutical compositions may be delivered
 15 by intranasal sprays, inhalation, and/or other aerosol delivery vehicles. Methods for delivering genes, nucleic acids, and peptide compositions directly to the lungs *via* nasal aerosol sprays has been described *e.g.*, in U. S. Patent 5,756,353 and U. S. Patent 5,804,212 (each specifically incorporated herein by reference in its entirety). Likewise, the delivery of drugs using intranasal microparticle resins (Takenaga *et al.*, 1998) and lysophosphatidyl-
 20 glycerol compounds (U. S. Patent 5,725,871, specifically incorporated herein by reference in its entirety) are also well-known in the pharmaceutical arts. Likewise, transmucosal drug delivery in the form of a polytetrafluoroethylene support matrix is described in U. S. Patent 5,780,045 (specifically incorporated herein by reference in its entirety).

4. LIPOSOME-, NANOCAPSULE-, AND MICROPARTICLE-MEDIATED DELIVERY

25 In certain embodiments, the inventors contemplate the use of liposomes, nanocapsules, microparticles, microspheres, lipid particles, vesicles, and the like, for the introduction of the compositions of the present invention into suitable host cells. In

particular, the compositions of the present invention may be formulated for delivery either encapsulated in a lipid particle, a liposome, a vesicle, a nanosphere, or a nanoparticle or the like.

Such formulations may be preferred for the introduction of
 5 pharmaceutically-acceptable formulations of the nucleic acids or constructs disclosed herein. The formation and use of liposomes is generally known to those of skill in the art (see for example, Couvreur *et al.*, 1977; Couvreur, 1988; Lasic, 1998; which describes the use of liposomes and nanocapsules in the targeted antibiotic therapy for intracellular bacterial infections and diseases). Recently, liposomes were developed with improved
 10 serum stability and circulation half-times (Gabizon and Papahadjopoulos, 1988; Allen and Choun, 1987; U. S. Patent 5,741,516, specifically incorporated herein by reference in its entirety). Further, various methods of liposome and liposome like preparations as potential drug carriers have been reviewed (Takakura, 1998; Chandran *et al.*, 1997; Margalit, 1995; U. S. Patent 5,567,434; U. S. Patent 5,552,157; U. S. Patent 5,565,213; U. S. Patent
 15 5,738,868 and U. S. Patent 5,795,587, each specifically incorporated herein by reference in its entirety).

Liposomes have been used successfully with a number of cell types that are normally resistant to transfection by other procedures including T cell suspensions, primary hepatocyte cultures and PC 12 cells (Renneisen *et al.*, 1990; Muller *et al.*, 1990). In
 20 addition, liposomes are free of the DNA length constraints that are typical of viral-based delivery systems. Liposomes have been used effectively to introduce genes, drugs (Heath and Martin, 1986; Heath *et al.*, 1986; Balazsovits *et al.*, 1989; Fresta and Puglisi, 1996), radiotherapeutic agents (Pikul *et al.*, 1987), enzymes (Imaizumi *et al.*, 1990a; Imaizumi
 25 *et al.*, 1990b), viruses (Faller and Baltimore, 1984), transcription factors and allosteric effectors (Nicolau and Gersonde, 1979) into a variety of cultured cell lines and animals. In addition, several successful clinical trials examining the effectiveness of liposome-mediated drug delivery have been completed (Lopez-Berestein *et al.*, 1985a; 1985b; Coune, 1988; Sculier *et al.*, 1988). Furthermore, several studies suggest that the use of

liposomes is not associated with autoimmune responses, toxicity or gonadal localization after systemic delivery (Mori and Fukatsu, 1992).

Liposomes are formed from phospholipids that are dispersed in an aqueous medium and spontaneously form multilamellar concentric bilayer vesicles (also termed
 5 multilamellar vesicles (MLVs). MLVs generally have diameters of from 25 nm to 4 μ m. Sonication of MLVs results in the formation of small unilamellar vesicles (SUVs) with diameters in the range of 200 to 500 Å, containing an aqueous solution in the core.

Liposomes bear resemblance to cellular membranes and are contemplated for use in connection with the present invention as carriers for the peptide compositions.
 10 They are widely suitable as both water- and lipid-soluble substances can be entrapped, *i.e.* in the aqueous spaces and within the bilayer itself, respectively. It is possible that the drug-bearing liposomes may even be employed for site-specific delivery of active agents by selectively modifying the liposomal formulation.

In addition to the teachings of Couvreur *et al.* (1977; 1988), the following
 15 information may be utilized in generating liposomal formulations. Phospholipids can form a variety of structures other than liposomes when dispersed in water, depending on the molar ratio of lipid to water. At low ratios the liposome is the preferred structure. The physical characteristics of liposomes depend on pH, ionic strength and the presence of divalent cations. Liposomes can show low permeability to ionic and polar substances, but
 20 at elevated temperatures undergo a phase transition which markedly alters their permeability. The phase transition involves a change from a closely packed, ordered structure, known as the gel state, to a loosely packed, less-ordered structure, known as the fluid state. This occurs at a characteristic phase-transition temperature and results in an increase in permeability to ions, sugars and drugs.

25 In addition to temperature, exposure to proteins can alter the permeability of liposomes. Certain soluble proteins, such as cytochrome c, bind, deform and penetrate the bilayer, thereby causing changes in permeability. Cholesterol inhibits this penetration of proteins, apparently by packing the phospholipids more tightly. It is contemplated that the

most useful liposome formations for antibiotic and inhibitor delivery will contain cholesterol.

The ability to trap solutes varies between different types of liposomes. For example, MLVs are moderately efficient at trapping solutes, but SUVs are extremely inefficient. SUVs offer the advantage of homogeneity and reproducibility in size distribution, however, and a compromise between size and trapping efficiency is offered by large unilamellar vesicles (LUVs). These are prepared by ether evaporation and are three to four times more efficient at solute entrapment than MLVs.

In addition to liposome characteristics, an important determinant in entrapping compounds is the physicochemical properties of the compound itself. Polar compounds are trapped in the aqueous spaces and nonpolar compounds bind to the lipid bilayer of the vesicle. Polar compounds are released through permeation or when the bilayer is broken, but nonpolar compounds remain affiliated with the bilayer unless it is disrupted by temperature or exposure to lipoproteins. Both types show maximum efflux rates at the phase transition temperature.

Liposomes interact with cells *via* four different mechanisms: endocytosis by phagocytic cells of the reticuloendothelial system such as macrophages and neutrophils; adsorption to the cell surface, either by nonspecific weak hydrophobic or electrostatic forces, or by specific interactions with cell-surface components; fusion with the plasma cell membrane by insertion of the lipid bilayer of the liposome into the plasma membrane, with simultaneous release of liposomal contents into the cytoplasm; and by transfer of liposomal lipids to cellular or subcellular membranes, or vice versa, without any association of the liposome contents. It often is difficult to determine which mechanism is operative and more than one may operate at the same time.

The fate and disposition of intravenously injected liposomes depend on their physical properties, such as size, fluidity, and surface charge. They may persist in tissues for h or days, depending on their composition, and half lives in the blood range from min to several h. Larger liposomes, such as MLVs and LUVs, are taken up rapidly by phagocytic cells of the reticuloendothelial system, but physiology of the circulatory system restrains

the exit of such large species at most sites. They can exit only in places where large openings or pores exist in the capillary endothelium, such as the sinusoids of the liver or spleen. Thus, these organs are the predominate site of uptake. On the other hand, SUVs show a broader tissue distribution but still are sequestered highly in the liver and spleen. In
 5 general, this *in vivo* behavior limits the potential targeting of liposomes to only those organs and tissues accessible to their large size. These include the blood, liver, spleen, bone marrow, and lymphoid organs.

Targeting is generally not a limitation in terms of the present invention. However, should specific targeting be desired, methods are available for this to be
 10 accomplished. Antibodies may be used to bind to the liposome surface and to direct the antibody and its drug contents to specific antigenic receptors located on a particular cell-type surface. Carbohydrate determinants (glycoprotein or glycolipid cell-surface components that play a role in cell-cell recognition, interaction and adhesion) may also be used as recognition sites as they have potential in directing liposomes to particular cell
 15 types. Mostly, it is contemplated that intravenous injection of liposomal preparations would be used, but other routes of administration are also conceivable.

Alternatively, the invention provides for pharmaceutically-acceptable nanocapsule formulations of the compositions of the present invention. Nanocapsules can generally entrap compounds in a stable and reproducible way (Henry-Michelland *et al.*,
 20 1987; Quintanar-Guerrero *et al.*, 1998; Douglas *et al.*, 1987). To avoid side effects due to intracellular polymeric overloading, such ultrafine particles (sized around 0.1 μm) should be designed using polymers able to be degraded *in vivo*. Biodegradable polyalkyl-cyanoacrylate nanoparticles that meet these requirements are contemplated for use in the present invention. Such particles may be easily made, as described (Couvreur *et al.*,
 25 1980; 1988; zur Muhlen *et al.*, 1998; Zambaux *et al.* 1998; Pinto-Alphandry *et al.*, 1995 and U. S. Patent 5,145,684, specifically incorporated herein by reference in its entirety).

IMMUNOGENIC COMPOSITIONS

In certain preferred embodiments of the present invention, immunogenic compositions, or vaccines, are provided. The immunogenic compositions will generally comprise one or more pharmaceutical compositions, such as those discussed above, in combination with an immunostimulant. An immunostimulant may be any substance that enhances or potentiates an immune response (antibody and/or cell-mediated) to an exogenous antigen. Examples of immunostimulants include adjuvants, biodegradable microspheres (*e.g.*, polylactic galactide) and liposomes (into which the compound is incorporated; *see e.g.*, Fullerton, U.S. Patent No. 4,235,877). Vaccine preparation is generally described in, for example, M.F. Powell and M.J. Newman, eds., "Vaccine Design (the subunit and adjuvant approach)," Plenum Press (NY, 1995). Pharmaceutical compositions and immunogenic compositions, or vaccines, within the scope of the present invention may also contain other compounds, which may be biologically active or inactive. For example, one or more immunogenic portions of other tumor antigens may be present, either incorporated into a fusion polypeptide or as a separate compound, within the composition.

Illustrative immunogenic compositions may contain DNA encoding one or more of the polypeptides as described above, such that the polypeptide is generated *in situ*. As noted above, the DNA may be present within any of a variety of delivery systems known to those of ordinary skill in the art, including nucleic acid expression systems, bacteria and viral expression systems. Numerous gene delivery techniques are well known in the art, such as those described by Rolland, *Crit. Rev. Therap. Drug Carrier Systems* 15:143-198, 1998, and references cited therein. Appropriate nucleic acid expression systems contain the necessary DNA sequences for expression in the patient (such as a suitable promoter and terminating signal). Bacterial delivery systems involve the administration of a bacterium (such as *Bacillus-Calmette-Guerrin*) that expresses an immunogenic portion of the polypeptide on its cell surface or secretes such an epitope. In a preferred embodiment, the DNA may be introduced using a viral expression system (*e.g.*, vaccinia or other pox virus, retrovirus, or adenovirus), which may involve the use of a non-

pathogenic (defective), replication competent virus. Suitable systems are disclosed, for example, in Fisher-Hoch *et al.*, *Proc. Natl. Acad. Sci. USA* 86:317-321, 1989; Flexner *et al.*, *Ann. N.Y. Acad. Sci.* 569:86-103, 1989; Flexner *et al.*, *Vaccine* 8:17-21, 1990; U.S. Patent Nos. 4,603,112, 4,769,330, and 5,017,487; WO 89/01973; U.S. Patent
 5 No. 4,777,127; GB 2,200,651; EP 0,345,242; WO 91/02805; Berkner, *Biotechniques* 6:616-627, 1988; Rosenfeld *et al.*, *Science* 252:431-434, 1991; Kolls *et al.*, *Proc. Natl. Acad. Sci. USA* 91:215-219, 1994; Kass-Eisler *et al.*, *Proc. Natl. Acad. Sci. USA* 90:11498-11502, 1993; Guzman *et al.*, *Circulation* 88:2838-2848, 1993; and Guzman *et al.*, *Cir. Res.* 73:1202-1207, 1993. Techniques for incorporating DNA into such expression
 10 systems are well known to those of ordinary skill in the art. The DNA may also be "naked," as described, for example, in Ulmer *et al.*, *Science* 259:1745-1749, 1993 and reviewed by Cohen, *Science* 259:1691-1692, 1993. The uptake of naked DNA may be increased by coating the DNA onto biodegradable beads, which are efficiently transported into the cells. It will be apparent that an immunogenic composition may comprise both a
 15 polynucleotide and a polypeptide component. Such immunogenic compositions may provide for an enhanced immune response.

It will be apparent that an immunogenic composition may contain pharmaceutically acceptable salts of the polynucleotides and polypeptides provided herein. Such salts may be prepared from pharmaceutically acceptable non-toxic bases, including
 20 organic bases (*e.g.*, salts of primary, secondary and tertiary amines and basic amino acids) and inorganic bases (*e.g.*, sodium, potassium, lithium, ammonium, calcium and magnesium salts).

While any suitable carrier known to those of ordinary skill in the art may be employed in the immunogenic compositions of this invention, the type of carrier will vary
 25 depending on the mode of administration. Compositions of the present invention may be formulated for any appropriate manner of administration, including for example, topical, oral, nasal, intravenous, intracranial, intraperitoneal, subcutaneous or intramuscular administration. For parenteral administration, such as subcutaneous injection, the carrier preferably comprises water, saline, alcohol, a fat, a wax or a buffer. For oral

administration, any of the above carriers or a solid carrier, such as mannitol, lactose, starch, magnesium stearate, sodium saccharine, talcum, cellulose, glucose, sucrose, and magnesium carbonate, may be employed. Biodegradable microspheres (*e.g.*, polylactate polyglycolate) may also be employed as carriers for the pharmaceutical compositions of this invention. Suitable biodegradable microspheres are disclosed, for example, in U.S. Patent Nos. 4,897,268; 5,075,109; 5,928,647; 5,811,128; 5,820,883; 5,853,763; 5,814,344 and 5,942,252. One may also employ a carrier comprising the particulate-protein complexes described in U.S. Patent No. 5,928,647, which are capable of inducing a class I-restricted cytotoxic T lymphocyte responses in a host.

Such compositions may also comprise buffers (*e.g.*, neutral buffered saline or phosphate buffered saline), carbohydrates (*e.g.*, glucose, mannose, sucrose or dextrans), mannitol, proteins, polypeptides or amino acids such as glycine, antioxidants, bacteriostats, chelating agents such as EDTA or glutathione, adjuvants (*e.g.*, aluminum hydroxide), solutes that render the formulation isotonic, hypotonic or weakly hypertonic with the blood of a recipient, suspending agents, thickening agents and/or preservatives. Alternatively, compositions of the present invention may be formulated as a lyophilizate. Compounds may also be encapsulated within liposomes using well known technology.

Any of a variety of immunostimulants may be employed in the immunogenic compositions of this invention. For example, an adjuvant may be included. Most adjuvants contain a substance designed to protect the antigen from rapid catabolism, such as aluminum hydroxide or mineral oil, and a stimulator of immune responses, such as lipid A, *Bordetella pertussis* or *Mycobacterium tuberculosis* derived proteins. Suitable adjuvants are commercially available as, for example, Freund's Incomplete Adjuvant and Complete Adjuvant (Difco Laboratories, Detroit, MI); Merck Adjuvant 65 (Merck and Company, Inc., Rahway, NJ); AS-2 (SmithKline Beecham, Philadelphia, PA); aluminum salts such as aluminum hydroxide gel (alum) or aluminum phosphate; salts of calcium, iron or zinc; an insoluble suspension of acylated tyrosine; acylated sugars; cationically or anionically derivatized polysaccharides; polyphosphazenes; biodegradable microspheres;

monophosphoryl lipid A and quil A. Cytokines, such as GM-CSF or interleukin-2, -7, or -12, may also be used as adjuvants.

Within the immunogenic compositions provided herein, the adjuvant composition is preferably designed to induce an immune response predominantly of the Th1 type. High levels of Th1-type cytokines (*e.g.*, IFN- γ , TNF α , IL-2 and IL-12) tend to favor the induction of cell mediated immune responses to an administered antigen. In contrast, high levels of Th2-type cytokines (*e.g.*, IL-4, IL-5, IL-6 and IL-10) tend to favor the induction of humoral immune responses. Following application of an immunogenic composition as provided herein, a patient will support an immune response that includes Th1- and Th2-type responses. Within a preferred embodiment, in which a response is predominantly Th1-type, the level of Th1-type cytokines will increase to a greater extent than the level of Th2-type cytokines. The levels of these cytokines may be readily assessed using standard assays. For a review of the families of cytokines, see Mosmann and Coffman, *Ann. Rev. Immunol.* 7:145-173, 1989.

Preferred adjuvants for use in eliciting a predominantly Th1-type response include, for example, a combination of monophosphoryl lipid A, preferably 3-de-O-acylated monophosphoryl lipid A (3D-MPL), together with an aluminum salt. MPL adjuvants are available from Corixa Corporation (Seattle, WA; *see* US Patent Nos. 4,436,727; 4,877,611; 4,866,034 and 4,912,094). CpG-containing oligonucleotides (in which the CpG dinucleotide is unmethylated) also induce a predominantly Th1 response. Such oligonucleotides are well known and are described, for example, in WO 96/02555, WO 99/33488 and U.S. Patent Nos. 6,008,200 and 5,856,462. Immunostimulatory DNA sequences are also described, for example, by Sato *et al.*, *Science* 273:352, 1996. Another preferred adjuvant is a saponin, preferably QS21 (Aquila Biopharmaceuticals Inc., Framingham, MA), which may be used alone or in combination with other adjuvants. For example, an enhanced system involves the combination of a monophosphoryl lipid A and saponin derivative, such as the combination of QS21 and 3D-MPL as described in WO 94/00153, or a less reactogenic composition where the QS21 is quenched with cholesterol, as described in WO 96/33739. Other preferred formulations comprise an oil-in-water

emulsion and tocopherol. A particularly potent adjuvant formulation involving QS21, 3D-MPL and tocopherol in an oil-in-water emulsion is described in WO 95/17210.

Other preferred adjuvants include Montanide ISA 720 (Seppic, France), SAF (Chiron, California, United States), ISCOMS (CSL), MF-59 (Chiron), the SBAS series of adjuvants (*e.g.*, SBAS-2 or SBAS-4, available from SmithKline Beecham, Rixensart, Belgium), Detox (Corixa, Hamilton, MT), RC-529 (Corixa, Hamilton, MT) and other aminoalkyl glucosaminide 4-phosphates (AGPs), such as those described in pending U.S. Patent Application Serial Nos. 08/853,826 and 09/074,720, the disclosures of which are incorporated herein by reference in their entireties.

Any immunogenic composition provided herein may be prepared using well known methods that result in a combination of antigen, immune response enhancer and a suitable carrier or excipient. The compositions described herein may be administered as part of a sustained release formulation (*i.e.*, a formulation such as a capsule, sponge or gel (composed of polysaccharides, for example) that effects a slow release of compound following administration). Such formulations may generally be prepared using well known technology (*see, e.g.*, Coombes *et al.*, *Vaccine* 14:1429-1438, 1996) and administered by, for example, oral, rectal or subcutaneous implantation, or by implantation at the desired target site. Sustained-release formulations may contain a polypeptide, polynucleotide or antibody dispersed in a carrier matrix and/or contained within a reservoir surrounded by a rate controlling membrane.

Carriers for use within such formulations are biocompatible, and may also be biodegradable; preferably the formulation provides a relatively constant level of active component release. Such carriers include microparticles of poly(lactide-co-glycolide), polyacrylate, latex, starch, cellulose, dextran and the like. Other delayed-release carriers include supramolecular biovectors, which comprise a non-liquid hydrophilic core (*e.g.*, a cross-linked polysaccharide or oligosaccharide) and, optionally, an external layer comprising an amphiphilic compound, such as a phospholipid (*see e.g.*, U.S. Patent No. 5,151,254 and PCT applications WO 94/20078, WO/94/23701 and WO 96/06638). The amount of active compound contained within a sustained release formulation depends upon

the site of implantation, the rate and expected duration of release and the nature of the condition to be treated or prevented.

Any of a variety of delivery vehicles may be employed within pharmaceutical compositions and immunogenic compositions to facilitate production of an antigen-specific immune response that targets tumor cells. Delivery vehicles include antigen presenting cells (APCs), such as dendritic cells, macrophages, B cells, monocytes and other cells that may be engineered to be efficient APCs. Such cells may, but need not, be genetically modified to increase the capacity for presenting the antigen, to improve activation and/or maintenance of the T cell response, to have anti-tumor effects *per se* and/or to be immunologically compatible with the receiver (*i.e.*, matched HLA haplotype). APCs may generally be isolated from any of a variety of biological fluids and organs, including tumor and peritumoral tissues, and may be autologous, allogeneic, syngeneic or xenogeneic cells.

Certain preferred embodiments of the present invention use dendritic cells or progenitors thereof as antigen-presenting cells. Dendritic cells are highly potent APCs (Banchereau and Steinman, *Nature* 392:245-251, 1998) and have been shown to be effective as a physiological adjuvant for eliciting prophylactic or therapeutic antitumor immunity (*see* Timmerman and Levy, *Ann. Rev. Med.* 50:507-529, 1999). In general, dendritic cells may be identified based on their typical shape (stellate *in situ*, with marked cytoplasmic processes (dendrites) visible *in vitro*), their ability to take up, process and present antigens with high efficiency and their ability to activate naïve T cell responses. Dendritic cells may, of course, be engineered to express specific cell-surface receptors or ligands that are not commonly found on dendritic cells *in vivo* or *ex vivo*, and such modified dendritic cells are contemplated by the present invention. As an alternative to dendritic cells, secreted vesicles antigen-loaded dendritic cells (called exosomes) may be used within a vaccine, or immunogenic composition (*see* Zitvogel *et al.*, *Nature Med.* 4:594-600, 1998).

Dendritic cells and progenitors may be obtained from peripheral blood, bone marrow, tumor-infiltrating cells, peritumoral tissues-infiltrating cells, lymph nodes, spleen,

skin, umbilical cord blood or any other suitable tissue or fluid. For example, dendritic cells may be differentiated *ex vivo* by adding a combination of cytokines such as GM-CSF, IL-4, IL-13 and/or TNF α to cultures of monocytes harvested from peripheral blood. Alternatively, CD34 positive cells harvested from peripheral blood, umbilical cord blood or
 5 bone marrow may be differentiated into dendritic cells by adding to the culture medium combinations of GM-CSF, IL-3, TNF α , CD40 ligand, LPS, flt3 ligand and/or other compound(s) that induce differentiation, maturation and proliferation of dendritic cells.

Dendritic cells are conveniently categorized as "immature" and "mature" cells, which allows a simple way to discriminate between two well characterized
 10 phenotypes. However, this nomenclature should not be construed to exclude all possible intermediate stages of differentiation. Immature dendritic cells are characterized as APC with a high capacity for antigen uptake and processing, which correlates with the high expression of Fc γ receptor and mannose receptor. The mature phenotype is typically characterized by a lower expression of these markers, but a high expression of cell surface
 15 molecules responsible for T cell activation such as class I and class II MHC, adhesion molecules (*e.g.*, CD54 and CD11) and costimulatory molecules (*e.g.*, CD40, CD80, CD86 and 4-1BB).

APCs may generally be transfected with a polynucleotide encoding a lung tumor protein (or portion or other variant thereof) such that the lung tumor polypeptide, or
 20 an immunogenic portion thereof, is expressed on the cell surface. Such transfection may take place *ex vivo*, and a composition comprising such transfected cells may then be used for therapeutic purposes, as described herein. Alternatively, a gene delivery vehicle that targets a dendritic or other antigen presenting cell may be administered to a patient, resulting in transfection that occurs *in vivo*. *In vivo* and *ex vivo* transfection of dendritic
 25 cells, for example, may generally be performed using any methods known in the art, such as those described in WO 97/24447, or the gene gun approach described by Mahvi *et al.*, *Immunology and cell Biology* 75:456-460, 1997. Antigen loading of dendritic cells may be achieved by incubating dendritic cells or progenitor cells with the lung tumor polypeptide, DNA (naked or within a plasmid vector) or RNA; or with antigen-expressing recombinant

bacterium or viruses (*e.g.*, vaccinia, fowlpox, adenovirus or lentivirus vectors). Prior to loading, the polypeptide may be covalently conjugated to an immunological partner that provides T cell help (*e.g.*, a carrier molecule). Alternatively, a dendritic cell may be pulsed with a non-conjugated immunological partner, separately or in the presence of the polypeptide.

Immunogenic compositions and pharmaceutical compositions may be presented in unit-dose or multi-dose containers, such as sealed ampoules or vials. Such containers are preferably hermetically sealed to preserve sterility of the formulation until use. In general, formulations may be stored as suspensions, solutions or emulsions in oily or aqueous vehicles. Alternatively, an immunogenic or pharmaceutical composition may be stored in a freeze-dried condition requiring only the addition of a sterile liquid carrier immediately prior to use.

CANCER THERAPY

In further aspects of the present invention, the compositions described herein may be used for immunotherapy of cancer, such as lung cancer. Within such methods, compositions are typically administered to a patient. As used herein, a "patient" refers to any warm-blooded animal, preferably a human. A patient may or may not be afflicted with cancer. Accordingly, the above pharmaceutical compositions and immunogenic compositions may be used to prevent the development of a cancer or to treat a patient afflicted with a cancer. A cancer may be diagnosed using criteria generally accepted in the art, including the presence of a malignant tumor. Pharmaceutical compositions and immunogenic compositions may be administered either prior to or following surgical removal of primary tumors and/or treatment such as administration of radiotherapy or conventional chemotherapeutic drugs. Administration may be by any suitable method, including administration by intravenous, intraperitoneal, intramuscular, subcutaneous, intranasal, intradermal, anal, vaginal, topical and oral routes.

Within certain embodiments, immunotherapy may be active immunotherapy, in which treatment relies on the *in vivo* stimulation of the endogenous host

immune system to react against tumors with the administration of immune response-modifying agents (such as polypeptides and polynucleotides as provided herein).

Within other embodiments, immunotherapy may be passive immunotherapy, in which treatment involves the delivery of agents with established tumor-immune reactivity (such as effector cells or antibodies) that can directly or indirectly mediate antitumor effects and does not necessarily depend on an intact host immune system. Examples of effector cells include T cells as discussed above, T lymphocytes (such as CD8⁺ cytotoxic T lymphocytes and CD4⁺ T-helper tumor-infiltrating lymphocytes), killer cells (such as Natural Killer cells and lymphokine-activated killer cells), B cells and antigen-presenting cells (such as dendritic cells and macrophages) expressing a polypeptide provided herein. T cell receptors and antibody receptors specific for the polypeptides recited herein may be cloned, expressed and transferred into other vectors or effector cells for adoptive immunotherapy. The polypeptides provided herein may also be used to generate antibodies or anti-idiotypic antibodies (as described above and in U.S. Patent No. 4,918,164) for passive immunotherapy.

Effector cells may generally be obtained in sufficient quantities for adoptive immunotherapy by growth *in vitro*, as described herein. Culture conditions for expanding single antigen-specific effector cells to several billion in number with retention of antigen recognition *in vivo* are well known in the art. Such *in vitro* culture conditions typically use intermittent stimulation with antigen, often in the presence of cytokines (such as IL-2) and non-dividing feeder cells. As noted above, immunoreactive polypeptides as provided herein may be used to rapidly expand antigen-specific T cell cultures in order to generate a sufficient number of cells for immunotherapy. In particular, antigen-presenting cells, such as dendritic, macrophage, monocyte, fibroblast and/or B cells, may be pulsed with immunoreactive polypeptides or transfected with one or more polynucleotides using standard techniques well known in the art. For example, antigen-presenting cells can be transfected with a polynucleotide having a promoter appropriate for increasing expression in a recombinant virus or other expression system. Cultured effector cells for use in therapy must be able to grow and distribute widely, and to survive long term *in vivo*.

Studies have shown that cultured effector cells can be induced to grow *in vivo* and to survive long term in substantial numbers by repeated stimulation with antigen supplemented with IL-2 (*see, for example, Cheever et al., Immunological Reviews 157:177, 1997*).

5 Alternatively, a vector expressing a polypeptide recited herein may be introduced into antigen presenting cells taken from a patient and clonally propagated *ex vivo* for transplant back into the same patient. Transfected cells may be reintroduced into the patient using any means known in the art, preferably in sterile form by intravenous, intracavitary, intraperitoneal or intratumor administration.

10 Routes and frequency of administration of the therapeutic compositions described herein, as well as dosage, will vary from individual to individual, and may be readily established using standard techniques. In general, the pharmaceutical compositions and immunogenic compositions may be administered by injection (*e.g., intracutaneous, intramuscular, intravenous or subcutaneous*), intranasally (*e.g., by aspiration*) or orally.

15 Preferably, between 1 and 10 doses may be administered over a 52 week period. Preferably, 6 doses are administered, at intervals of 1 month, and booster vaccinations may be given periodically thereafter. Alternate protocols may be appropriate for individual patients. A suitable dose is an amount of a compound that, when administered as described above, is capable of promoting an anti-tumor immune response, and is at least 10-50%

20 above the basal (*i.e., untreated*) level. Such response can be monitored by measuring the anti-tumor antibodies in a patient or by vaccine-dependent generation of cytolytic effector cells capable of killing the patient's tumor cells *in vitro*. Such vaccines, or immunogenic compositions, should also be capable of causing an immune response that leads to an improved clinical outcome (*e.g., more frequent remissions, complete or partial or longer*

25 disease-free survival) in vaccinated patients as compared to non-vaccinated patients. In general, for compositions comprising one or more polypeptides, the amount of each polypeptide present in a dose ranges from about 25 μ g to 5 mg per kg of host. Suitable dose sizes will vary with the size of the patient, but will typically range from about 0.1 mL to about 5 mL.

In general, an appropriate dosage and treatment regimen provides the active compound(s) in an amount sufficient to provide therapeutic and/or prophylactic benefit. Such a response can be monitored by establishing an improved clinical outcome (*e.g.*, more frequent remissions, complete or partial, or longer disease-free survival) in treated patients as compared to non-treated patients. Increases in preexisting immune responses to a lung tumor protein generally correlate with an improved clinical outcome. Such immune responses may generally be evaluated using standard proliferation, cytotoxicity or cytokine assays, which may be performed using samples obtained from a patient before and after treatment.

10 **CANCER DETECTION AND DIAGNOSIS**

In general, a cancer may be detected in a patient based on the presence of one or more lung tumor proteins and/or polynucleotides encoding such proteins in a biological sample (for example, blood, sera, sputum urine and/or tumor biopsies) obtained from the patient. In other words, such proteins may be used as markers to indicate the presence or absence of a cancer such as lung cancer. In addition, such proteins may be useful for the detection of other cancers. The binding agents provided herein generally permit detection of the level of antigen that binds to the agent in the biological sample. Polynucleotide primers and probes may be used to detect the level of mRNA encoding a tumor protein, which is also indicative of the presence or absence of a cancer. In general, a lung tumor sequence should be present at a level that is at least three fold higher in tumor tissue than in normal tissue

There are a variety of assay formats known to those of ordinary skill in the art for using a binding agent to detect polypeptide markers in a sample. *See, e.g.*, Harlow and Lane, *Antibodies: A Laboratory Manual*, Cold Spring Harbor Laboratory, 1988. In general, the presence or absence of a cancer in a patient may be determined by (a) contacting a biological sample obtained from a patient with a binding agent; (b) detecting in the sample a level of polypeptide that binds to the binding agent; and (c) comparing the level of polypeptide with a predetermined cut-off value.

In a preferred embodiment, the assay involves the use of binding agent immobilized on a solid support to bind to and remove the polypeptide from the remainder of the sample. The bound polypeptide may then be detected using a detection reagent that contains a reporter group and specifically binds to the binding agent/polypeptide complex.

5 Such detection reagents may comprise, for example, a binding agent that specifically binds to the polypeptide or an antibody or other agent that specifically binds to the binding agent, such as an anti-immunoglobulin, protein G, protein A or a lectin. Alternatively, a competitive assay may be utilized, in which a polypeptide is labeled with a reporter group and allowed to bind to the immobilized binding agent after incubation of the binding agent

10 with the sample. The extent to which components of the sample inhibit the binding of the labeled polypeptide to the binding agent is indicative of the reactivity of the sample with the immobilized binding agent. Suitable polypeptides for use within such assays include full length lung tumor proteins and portions thereof to which the binding agent binds, as described above.

15 The solid support may be any material known to those of ordinary skill in the art to which the tumor protein may be attached. For example, the solid support may be a test well in a microtiter plate or a nitrocellulose or other suitable membrane. Alternatively, the support may be a bead or disc, such as glass, fiberglass, latex or a plastic material such as polystyrene or polyvinylchloride. The support may also be a magnetic

20 particle or a fiber optic sensor, such as those disclosed, for example, in U.S. Patent No. 5,359,681. The binding agent may be immobilized on the solid support using a variety of techniques known to those of skill in the art, which are amply described in the patent and scientific literature. In the context of the present invention, the term "immobilization" refers to both noncovalent association, such as adsorption, and covalent attachment (which

25 may be a direct linkage between the agent and functional groups on the support or may be a linkage by way of a cross-linking agent). Immobilization by adsorption to a well in a microtiter plate or to a membrane is preferred. In such cases, adsorption may be achieved by contacting the binding agent, in a suitable buffer, with the solid support for a suitable amount of time. The contact time varies with temperature, but is typically between about 1

hour and about 1 day. In general, contacting a well of a plastic microtiter plate (such as polystyrene or polyvinylchloride) with an amount of binding agent ranging from about 10 ng to about 10 μ g, and preferably about 100 ng to about 1 μ g, is sufficient to immobilize an adequate amount of binding agent.

5 Covalent attachment of binding agent to a solid support may generally be achieved by first reacting the support with a bifunctional reagent that will react with both the support and a functional group, such as a hydroxyl or amino group, on the binding agent. For example, the binding agent may be covalently attached to supports having an appropriate polymer coating using benzoquinone or by condensation of an aldehyde group
10 on the support with an amine and an active hydrogen on the binding partner (*see, e.g.*, Pierce Immunotechnology Catalog and Handbook, 1991, at A12-A13).

In certain embodiments, the assay is a two-antibody sandwich assay. This assay may be performed by first contacting an antibody that has been immobilized on a solid support, commonly the well of a microtiter plate, with the sample, such that
15 polypeptides within the sample are allowed to bind to the immobilized antibody. Unbound sample is then removed from the immobilized polypeptide-antibody complexes and a detection reagent (preferably a second antibody capable of binding to a different site on the polypeptide) containing a reporter group is added. The amount of detection reagent that remains bound to the solid support is then determined using a method appropriate for the
20 specific reporter group.

More specifically, once the antibody is immobilized on the support as described above, the remaining protein binding sites on the support are typically blocked. Any suitable blocking agent known to those of ordinary skill in the art, such as bovine serum albumin or Tween 20TM (Sigma Chemical Co., St. Louis, MO). The immobilized
25 antibody is then incubated with the sample, and polypeptide is allowed to bind to the antibody. The sample may be diluted with a suitable diluent, such as phosphate-buffered saline (PBS) prior to incubation. In general, an appropriate contact time (*i.e.*, incubation time) is a period of time that is sufficient to detect the presence of polypeptide within a sample obtained from an individual with lung cancer. Preferably, the contact time is

sufficient to achieve a level of binding that is at least about 95% of that achieved at equilibrium between bound and unbound polypeptide. Those of ordinary skill in the art will recognize that the time necessary to achieve equilibrium may be readily determined by assaying the level of binding that occurs over a period of time. At room temperature, an incubation time of about 30 minutes is generally sufficient.

Unbound sample may then be removed by washing the solid support with an appropriate buffer, such as PBS containing 0.1% Tween 20™. The second antibody, which contains a reporter group, may then be added to the solid support. Preferred reporter groups include those groups recited above.

The detection reagent is then incubated with the immobilized antibody-polypeptide complex for an amount of time sufficient to detect the bound polypeptide. An appropriate amount of time may generally be determined by assaying the level of binding that occurs over a period of time. Unbound detection reagent is then removed and bound detection reagent is detected using the reporter group. The method employed for detecting the reporter group depends upon the nature of the reporter group. For radioactive groups, scintillation counting or autoradiographic methods are generally appropriate. Spectroscopic methods may be used to detect dyes, luminescent groups and fluorescent groups. Biotin may be detected using avidin, coupled to a different reporter group (commonly a radioactive or fluorescent group or an enzyme). Enzyme reporter groups may generally be detected by the addition of substrate (generally for a specific period of time), followed by spectroscopic or other analysis of the reaction products.

To determine the presence or absence of a cancer, such as lung cancer, the signal detected from the reporter group that remains bound to the solid support is generally compared to a signal that corresponds to a predetermined cut-off value. In one preferred embodiment, the cut-off value for the detection of a cancer is the average mean signal obtained when the immobilized antibody is incubated with samples from patients without the cancer. In general, a sample generating a signal that is three standard deviations above the predetermined cut-off value is considered positive for the cancer. In an alternate preferred embodiment, the cut-off value is determined using a Receiver Operator Curve,

according to the method of Sackett *et al.*, *Clinical Epidemiology: A Basic Science for Clinical Medicine*, Little Brown and Co., 1985, p. 106-7. Briefly, in this embodiment, the cut-off value may be determined from a plot of pairs of true positive rates (*i.e.*, sensitivity) and false positive rates (100%-specificity) that correspond to each possible cut-off value
 5 for the diagnostic test result. The cut-off value on the plot that is the closest to the upper left-hand corner (*i.e.*, the value that encloses the largest area) is the most accurate cut-off value, and a sample generating a signal that is higher than the cut-off value determined by this method may be considered positive. Alternatively, the cut-off value may be shifted to the left along the plot, to minimize the false positive rate, or to the right, to minimize the
 10 false negative rate. In general, a sample generating a signal that is higher than the cut-off value determined by this method is considered positive for a cancer.

In a related embodiment, the assay is performed in a flow-through or strip test format, wherein the binding agent is immobilized on a membrane, such as nitrocellulose. In the flow-through test, polypeptides within the sample bind to the
 15 immobilized binding agent as the sample passes through the membrane. A second, labeled binding agent then binds to the binding agent-polypeptide complex as a solution containing the second binding agent flows through the membrane. The detection of bound second binding agent may then be performed as described above. In the strip test format, one end of the membrane to which binding agent is bound is immersed in a solution containing the
 20 sample. The sample migrates along the membrane through a region containing second binding agent and to the area of immobilized binding agent. Concentration of second binding agent at the area of immobilized antibody indicates the presence of a cancer. Typically, the concentration of second binding agent at that site generates a pattern, such as a line, that can be read visually. The absence of such a pattern indicates a negative result.
 25 In general, the amount of binding agent immobilized on the membrane is selected to generate a visually discernible pattern when the biological sample contains a level of polypeptide that would be sufficient to generate a positive signal in the two-antibody sandwich assay, in the format discussed above. Preferred binding agents for use in such assays are antibodies and antigen-binding fragments thereof. Preferably, the amount of

antibody immobilized on the membrane ranges from about 25 ng to about 1 μ g, and more preferably from about 50 ng to about 500 ng. Such tests can typically be performed with a very small amount of biological sample.

Of course, numerous other assay protocols exist that are suitable for use
 5 with the tumor proteins or binding agents of the present invention. The above descriptions are intended to be exemplary only. For example, it will be apparent to those of ordinary skill in the art that the above protocols may be readily modified to use lung tumor polypeptides to detect antibodies that bind to such polypeptides in a biological sample. The detection of such lung tumor protein specific antibodies may correlate with the
 10 presence of a cancer.

A cancer may also, or alternatively, be detected based on the presence of T cells that specifically react with a lung tumor protein in a biological sample. Within certain methods, a biological sample comprising CD4⁺ and/or CD8⁺ T cells isolated from a patient is incubated with a lung tumor polypeptide, a polynucleotide encoding such a polypeptide
 15 and/or an APC that expresses at least an immunogenic portion of such a polypeptide, and the presence or absence of specific activation of the T cells is detected. Suitable biological samples include, but are not limited to, isolated T cells. For example, T cells may be isolated from a patient by routine techniques (such as by Ficoll/Hypaque density gradient centrifugation of peripheral blood lymphocytes). T cells may be incubated *in vitro* for 2-9
 20 days (typically 4 days) at 37°C with polypeptide (*e.g.*, 5 - 25 μ g/ml). It may be desirable to incubate another aliquot of a T cell sample in the absence of lung tumor polypeptide to serve as a control. For CD4⁺ T cells, activation is preferably detected by evaluating proliferation of the T cells. For CD8⁺ T cells, activation is preferably detected by evaluating cytolytic activity. A level of proliferation that is at least two fold greater and/or
 25 a level of cytolytic activity that is at least 20% greater than in disease-free patients indicates the presence of a cancer in the patient.

As noted above, a cancer may also, or alternatively, be detected based on the level of mRNA encoding a lung tumor protein in a biological sample. For example, at least two oligonucleotide primers may be employed in a polymerase chain reaction (PCR) based

assay to amplify a portion of a lung tumor cDNA derived from a biological sample, wherein at least one of the oligonucleotide primers is specific for (*i.e.*, hybridizes to) a polynucleotide encoding the lung tumor protein. The amplified cDNA is then separated and detected using techniques well known in the art, such as gel electrophoresis. Similarly,
 5 oligonucleotide probes that specifically hybridize to a polynucleotide encoding a lung tumor protein may be used in a hybridization assay to detect the presence of polynucleotide encoding the tumor protein in a biological sample.

To permit hybridization under assay conditions, oligonucleotide primers and probes should comprise an oligonucleotide sequence that has at least about 60%, preferably
 10 at least about 75% and more preferably at least about 90%, identity to a portion of a polynucleotide encoding a lung tumor protein that is at least 10 nucleotides, and preferably at least 20 nucleotides, in length. Preferably, oligonucleotide primers and/or probes hybridize to a polynucleotide encoding a polypeptide described herein under moderately stringent conditions, as defined above. Oligonucleotide primers and/or probes which may
 15 be usefully employed in the diagnostic methods described herein preferably are at least 10-40 nucleotides in length. In a preferred embodiment, the oligonucleotide primers comprise at least 10 contiguous nucleotides, more preferably at least 15 contiguous nucleotides, of a DNA molecule having a sequence recited in SEQ ID NO: 1, 11-13, 15, 20, 23-27, 29, 30, 33, 34, 39, 41, 43-46, 51, 52, 57, 58, 60, 62, 65-67, 69-71, 74, 76, 79, 80, 84, 86, 89-92, 95,
 20 97, 98, 101, 110, 111, 113-119, 121-128, 130-134, 136, 138, 139, 141, 143, 146-151, 153, 154, 157-160, 162-164, 167-178, 180, 181, 183, 186-190, 192, 193, 195-220, 224, 226-231, 234, 236, 237, 240, 241, 244-246, 248, 254, 255, 261, 262, 266, 270, 275, 280, 282, 283, 288, 289, 290, 292, 295, 301, 303, 304, 309, 311, 341-782, 784, 785, 790, 792, 794, 796, 800-804, 807, 808, 810-826, 828-1664, 1669 or 1676. Techniques for both PCR
 25 based assays and hybridization assays are well known in the art (*see*, for example, Mullis *et al.*, *Cold Spring Harbor Symp. Quant. Biol.*, 51:263, 1987; Erlich ed., *PCR Technology*, Stockton Press, NY, 1989).

One preferred assay employs RT-PCR, in which PCR is applied in conjunction with reverse transcription. Typically, RNA is extracted from a biological

sample, such as biopsy tissue, and is reverse transcribed to produce cDNA molecules. PCR amplification using at least one specific primer generates a cDNA molecule, which may be separated and visualized using, for example, gel electrophoresis. Amplification may be performed on biological samples taken from a test patient and from an individual who is not afflicted with a cancer. The amplification reaction may be performed on several dilutions of cDNA spanning two orders of magnitude. A two-fold or greater increase in expression in several dilutions of the test patient sample as compared to the same dilutions of the non-cancerous sample is typically considered positive.

In another embodiment, the compositions described herein may be used as markers for the progression of cancer. In this embodiment, assays as described above for the diagnosis of a cancer may be performed over time, and the change in the level of reactive polypeptide(s) or polynucleotide(s) evaluated. For example, the assays may be performed every 24-72 hours for a period of 6 months to 1 year, and thereafter performed as needed. In general, a cancer is progressing in those patients in whom the level of polypeptide or polynucleotide detected increases over time. In contrast, the cancer is not progressing when the level of reactive polypeptide or polynucleotide either remains constant or decreases with time.

Certain *in vivo* diagnostic assays may be performed directly on a tumor. One such assay involves contacting tumor cells with a binding agent. The bound binding agent may then be detected directly or indirectly via a reporter group. Such binding agents may also be used in histological applications. Alternatively, polynucleotide probes may be used within such applications.

As noted above, to improve sensitivity, multiple lung tumor protein markers may be assayed within a given sample. It will be apparent that binding agents specific for different proteins provided herein may be combined within a single assay. Further, multiple primers or probes may be used concurrently. The selection of tumor protein markers may be based on routine experiments to determine combinations that results in optimal sensitivity. In addition, or alternatively, assays for tumor proteins provided herein may be combined with assays for other known tumor antigens.

DIAGNOSTIC KITS

The present invention further provides kits for use within any of the above diagnostic methods. Such kits typically comprise two or more components necessary for performing a diagnostic assay. Components may be compounds, reagents, containers
5 and/or equipment. For example, one container within a kit may contain a monoclonal antibody or fragment thereof that specifically binds to a lung tumor protein. Such antibodies or fragments may be provided attached to a support material, as described above. One or more additional containers may enclose elements, such as reagents or
10 buffers, to be used in the assay. Such kits may also, or alternatively, contain a detection reagent as described above that contains a reporter group suitable for direct or indirect detection of antibody binding.

Alternatively, a kit may be designed to detect the level of mRNA encoding a lung tumor protein in a biological sample. Such kits generally comprise at least one oligonucleotide probe or primer, as described above, that hybridizes to a polynucleotide
15 encoding a lung tumor protein. Such an oligonucleotide may be used, for example, within a PCR or hybridization assay. Additional components that may be present within such kits include a second oligonucleotide and/or a diagnostic reagent or container to facilitate the detection of a polynucleotide encoding a lung tumor protein.

The following Examples are offered by way of illustration and not by way
20 of limitation.

EXAMPLE 1IDENTIFICATION AND CHARACTERIZATION OF LUNG
TUMOR PROTEIN cDNAS

5 This Example illustrates the identification of cDNA molecules encoding lung tumor proteins.

A. Isolation of cDNA Sequences from Lung Adenocarcinoma Libraries using Conventional cDNA Library Subtraction

10 A human lung adenocarcinoma cDNA expression library was constructed from poly A⁺ RNA from patient tissues (# 40031486) using a Superscript Plasmid System for cDNA Synthesis and Plasmid Cloning kit (BRL Life Technologies, Gaithersburg, MD) following the manufacturer's protocol. Specifically, lung carcinoma tissues were homogenized with polytron (Kinematica, Switzerland) and total RNA was extracted using Trizol reagent (BRL Life Technologies) as directed by the manufacturer. The poly A⁺ RNA was then purified using an oligo dT cellulose column as described in Sambrook et al.,
15 *Molecular Cloning: A Laboratory Manual*, Cold Spring Harbor Laboratories, Cold Spring Harbor, NY, 1989. First-strand cDNA was synthesized using the NotI/Oligo-dT18 primer. Double-stranded cDNA was synthesized, ligated with BstXI/EcoRI adaptors (Invitrogen, San Diego, CA) and digested with NotI. Following size fractionation with cDNA size
20 fractionation columns (BRL Life Technologies), the cDNA was ligated into the BstXI/NotI site of pcDNA3.1 (Invitrogen) and transformed into ElectroMax *E. coli* DH10B cells (BRL Life Technologies) by electroporation. A total of 3 x 10⁶ independent colonies were generated.

25 Using the same procedure, a normal human cDNA expression library was prepared from a panel of normal tissue specimens, including lung, liver, pancreas, skin, kidney, brain and resting PBMC.

 cDNA library subtraction was performed using the above lung adenocarcinoma and normal tissue cDNA libraries, as described by Hara *et al.* (*Blood*,

84:189-199, 1994) with some modifications. Specifically, a lung adenocarcinoma-specific subtracted cDNA library was generated as follows. The normal tissue cDNA library (80 µg) was digested with BamHI and XhoI, followed by a filling-in reaction with DNA polymerase Klenow fragment. After phenol-chloroform extraction and ethanol precipitation, the DNA was dissolved in 133 µl of H₂O, heat-denatured and mixed with 133 µl (133 µg) of Photoprobe biotin (Vector Laboratories, Burlingame, CA). As recommended by the manufacturer, the resulting mixture was irradiated with a 270 W sunlamp on ice for 20 minutes. Additional Photoprobe biotin (67 µl) was added and the biotinylation reaction was repeated. After extraction with butanol five times, the DNA was ethanol-precipitated and dissolved in 23 µl H₂O. The resulting DNA, plus other highly redundant cDNA clones that were frequently recovered in previous lung subtractions formed the driver DNA.

To form the tracer DNA, 10 µg lung adenocarcinoma cDNA library was digested with NotI and SpeI, phenol chloroform extracted and passed through Chroma spin-400 columns (Clontech, Palo Alto, CA). Typically, 5 µg of cDNA was recovered after the sizing column. Following ethanol precipitation, the tracer DNA was dissolved in 5 µl H₂O. Tracer DNA was mixed with 15 µl driver DNA and 20 µl of 2 x hybridization buffer (1.5 M NaCl/10 mM EDTA/50 mM HEPES pH 7.5/0.2% sodium dodecyl sulfate), overlaid with mineral oil, and heat-denatured completely. The sample was immediately transferred into a 68 °C water bath and incubated for 20 hours (long hybridization [LH]). The reaction mixture was then subjected to a streptavidin treatment followed by phenol/chloroform extraction. This process was repeated three more times. Subtracted DNA was precipitated, dissolved in 12 µl H₂O, mixed with 8 µl driver DNA and 20 µl of 2 x hybridization buffer, and subjected to a hybridization at 68°C for 2 hours (short hybridization [SH]). After removal of biotinylated double-stranded DNA, subtracted cDNA was ligated into NotI/SpeI site of chloramphenicol resistant pBCSK⁺ (Stratagene, La Jolla, CA) and transformed into ElectroMax *E. coli* DH10B cells by electroporation to generate a lung adenocarcinoma specific subtracted cDNA library, referred to as LAT-S1

Similarly, LAT-S2 was generated by including 23 genes that were over-expressed in the tracer as additional drivers.

A second human lung adenocarcinoma cDNA expression library was constructed using adenocarcinoma tissue from a second patient (# 86-66) and used to
5 prepare a second lung adenocarcinoma-specific subtracted cDNA library (referred to as LAT2-S2), as described above, using the same panel of normal tissues and the additional genes over-expressed in LAT-S1.

A third human metastatic lung adenocarcinoma library was constructed from a pool of two lung pleural effusions with lung and gastric adenocarcinoma origins.
10 The subtracted cDNA library, Mets-sub2 was generated as described above using the same panel of normal tissues. However, the Mets-sub3 subtracted library was constructed by including 51 additional genes as drivers. These 51 genes were recovered in Mets-sub2, representing over-expressed housekeeping genes in the testers. As a result, Mets-sub3 is more complexed and normalized.

A total of 16 cDNA fragments isolated from LAT-S1, 585 cDNA fragments isolated from LAT-S2, 568 cDNA clones from LAT2-S2, 15 cDNA clones from Mets-sub2 and 343 cDNA clones from Mets-sub3, described above, were colony PCR amplified and their mRNA expression levels in lung tumor, normal lung, and various other normal and tumor tissues were determined using microarray technology (Incyte, Palo Alto, CA).
15 Briefly, the PCR amplification products were dotted onto slides in an array format, with each product occupying a unique location in the array. mRNA was extracted from the tissue sample to be tested, reverse transcribed, and fluorescent-labeled cDNA probes were generated. The microarrays were probed with the labeled cDNA probes, the slides scanned and fluorescence intensity was measured. This intensity correlates with the hybridization
20 intensity. Seventy-three non-redundant cDNA clones, of which 42 were found to be unique, showed over-expression in lung tumors, with expression in normal tissues tested (lung, skin, lymph node, colon, liver, pancreas, breast, heart, bone marrow, large intestine, kidney, stomach, brain, small intestine, bladder and salivary gland) being either undetectable, or at significantly lower levels compared to lung adenocarcinoma tumors.

These clones were further characterized by DNA sequencing with a Perkin Elmer/Applied Biosystems Division Automated Sequencer Model 373A and/or Model 377 (Foster City, CA).

The sequences were compared to known sequences in the gene bank using the EMBL GenBank databases (release 96). No significant homologies were found to the sequence provided in SEQ ID NO: 67, with no apparent homology to previously identified expressed sequence tags (ESTs). The sequences of SEQ ID NO: 60, 62, 65, 66, 69-71, 74, 76, 79, 80, 84, 86, 89-92, 95, 97 and 98 were found to show some homology to previously identified expressed sequence tags (ESTs). The cDNA sequences of SEQ ID NO: 59, 61, 63, 64, 67, 68, 72, 73, 75, 77, 78, 81-83, 85, 87, 88, 93, 94, 96, 99 and 100 showed homology to previously identified genes. The full-length cDNA sequences for the clones of SEQ ID NO: 96 and 100 are provided in SEQ ID NO: 316 and 318, respectively. The amino acid sequences for the clones of SEQ ID NO: 59, 61, 63, 64, 68, 73, 82, 83, 94, 96 and 100 are provided in SEQ ID NO: 331, 328, 329, 332, 327, 333, 330, 326, 325, 324 and 335, respectively. A predicted amino acid sequence encoded by the sequence of SEQ ID NO: 69 (referred to as L552S) is provided in SEQ ID NO: 786.

Further studies led to the isolation of an extended cDNA sequence, and open reading frame, for L552S (SEQ ID NO: 790). The predicted amino acid sequence encoded by the cDNA sequence of SEQ ID NO: 790 is provided in SEQ ID NO: 791. The determined cDNA sequence of an isoform of L552S is provided in SEQ ID NO: 792, with the corresponding predicted amino acid sequence being provided in SEQ ID NO: 793. Subsequent studies led to the isolation of the full-length cDNA sequence of L552S (SEQ ID NO: 808). The corresponding amino acid sequence is provided in SEQ ID NO: 809. No homologies were found to the protein sequence of L552S. However, nucleotides 533-769 of the full-length cDNA sequence were found to show homology to a previously identified DNA sequence.

Full-length cloning efforts on L552S led to the isolation of three additional cDNA sequences (SEQ ID NO: 810-812) from a metastatic lung adenocarcinoma library. The sequence of SEQ ID NO: 810 was found to show some homology to previously

identified human DNA sequences. The sequence of SEQ ID NO: 811 was found to show some homology to a previously identified DNA sequence. The sequence of SEQ ID NO: 812 was found to show some homology to previously identified ESTs.

The gene of SEQ ID NO: 84 (referred to as L551S) was determined by real-time RT-PCR analysis to be over-expressed in 2/9 primary adenocarcinomas and to be expressed at lower levels in 2/2 metastatic adenocarcinomas and 1/2 squamous cell carcinomas. No expression was observed in normal tissues, with the exception of very low expression in normal stomach. Further studies on L551S led to the isolation of the 5' and 3' cDNA consensus sequences provided in SEQ ID NO: 801 and 802, respectively. The L551S 5' sequence was found to show some homology to the previously identified gene STY8 (cDNA sequence provided in SEQ ID NO: 803; corresponding amino acid sequence provided in SEQ ID NO: 805), which is a mitogen activated protein kinase phosphatase. However, no significant homologies were found to the 3' sequence of L551S. Subsequently, an extended cDNA sequence for L551S was isolated (SEQ ID NO: 804). The corresponding amino acid sequence is provided in SEQ ID NO: 806. Further studies led to the isolation of two independent full-length clones for L551S (referred to as 54298 and 54305). These two clones have five nucleotide differences compared to the STY8 DNA sequence. Two of these differences are single nucleotide polymorphisms which do not effect the encoded amino acid sequences. The other three nucleotide differences are consistent between the two L551S clones but lead to encoded amino acid sequences that are different from the STY8 protein sequence. The determined cDNA sequences for the L551S full-length clones 54305 and 54298 are provided in SEQ ID NO: 825 and 826, respectively, with the amino acid sequence for L551S being provided in SEQ ID NO: 827.

B. Isolation of cDNA Sequences from Lung Adenocarcinoma Libraries using PCR-Based cDNA Library Subtraction

cDNA clones from a PCR-based subtraction library, containing cDNA from a pool of two human lung primary adenocarcinomas subtracted against a pool of nine normal human tissue cDNAs including skin, colon, lung, esophagus, brain, kidney, spleen,

pancreas and liver, (Clontech, Palo Alto, CA) were derived and submitted to a first round of PCR amplification. This library (referred to as ALT-1) was subjected to a second round of PCR amplification, following the manufacturer's protocol. The expression levels of 760 cDNA clones in lung tumor, normal lung, and various other normal and tumor tissues, were examined using microarray technology as described above. A total of 118 clones, of which 55 were unique, were found to be over-expressed in lung tumor tissue, with expression in normal tissues tested (lung, skin, lymph node, colon, liver, pancreas, breast, heart, bone marrow, large intestine, kidney, stomach, brain, small intestine, bladder and salivary gland) being either undetectable, or at significantly lower levels. The sequences were compared to known sequences in the gene bank using the EMBL and GenBank databases (release 96). No significant homologies (including ESTs) were found to the sequence provided in SEQ ID NO: 44. The sequences of SEQ ID NO: 1, 11, 13, 15, 20, 23-27, 29, 30, 33, 34, 39, 41, 43, 45, 46, 51 and 57 were found to show some homology to previously identified expressed sequence tags (ESTs). The cDNA sequences of SEQ ID NO: 2-10, 12, 14, 16-19, 21, 22, 28, 31, 32, 35-38, 40, 42, 44, 47-50, 52-56 and 58 showed homology to previously identified genes. The full-length cDNA sequences for the clones of SEQ ID NO: 18, 22, 31, 35, 36 and 42 are provided in SEQ ID NO: 320, 319, 323, 321, 317, 321 and 322, respectively, with the corresponding amino acid sequences being provided in SEQ ID NO: 337, 336, 340, 338, 334, and 339, respectively.

Further studies led to the isolation of an extended cDNA sequence for the clone of SEQ ID NO: 33 (referred to as L801P). This extended cDNA sequence (provided in SEQ ID NO: 796), was found to contain three potential open reading frames (ORFs). The predicted amino acid sequences encoded by these three ORFs are provided in SEQ ID NO: 797-799, respectively. Additional full-length cloning efforts led to still further extended cDNA sequence for L801P, set forth in SEQ ID NO:1669, in addition to five potential open reading frames (ORFs 4-9; SEQ ID NOs: 1670-1675, respectively) encoded by the extended cDNA sequence. Moreover, L801P was mapped to chromosomal region 20p13 and a 137 amino acid ORF from this genomic region was identified that corresponds to ORF4 (SEQ ID NO: 1670), suggesting that this is likely an authentic ORF for L801P.

By microarray analysis, L801P was overexpressed by 2-fold or greater in the lung tumor probe groups compared to the normal tissue probe group (not shown). By real-time PCR analysis, greater than 50% of lung adenocarcinoma and greater than 30% of lung squamous cell carcinoma tumor samples tested had elevated L801P expression
 5 relative to normal lung tissue. Of those that displayed elevated L801P, the level of expression was greater than 10-fold higher than in normal lung tissue samples. Moreover, low or no expression of L801P was detected in an extensive panel of normal tissue RNAs.

We have also found that L801P expression is detected in a number of other tumor types, including breast, prostate, ovarian and colon tumors, and thus may have
 10 diagnostic and/or therapeutic utility in these cancer types as well.

In subsequent studies, a full-length cDNA sequence for the clone of SEQ ID NO: 44 (referred to as L844P) was isolated (provided in SEQ ID NO: 800). Comparison of this sequence with those in the public databases revealed that the 470 bases at the 5' end of the sequence show homology to the known gene dihydrodiol dehydrogenase, thus
 15 indicating that L844P is a novel transcript of the dihydrodiol dehydrogenase family having 2007 base pairs of previously unidentified 3' untranslated region.

The predicted amino acid sequence encoded by the sequence of SEQ ID NO: 46 (referred to as L840P) is provided in SEQ ID NO: 787. An extended cDNA sequence for L840P, which was determined to include an open reading frame, is provided
 20 in SEQ ID NO: 794. The predicted amino acid sequence encoded by the cDNA sequence of SEQ ID NO: 794 is provided in SEQ ID NO: 795. The full-length cDNA sequence for the clone of SEQ ID NO: 54 (referred to as L548S) is provided in SEQ ID NO: 788, with the corresponding amino acid sequence being provided in SEQ ID NO: 789.

Northern blot analyses of the genes of SEQ ID NO: 25 and 46 (referred to as
 25 L839P and L840P, respectively) were remarkably similar. Both genes were expressed in 1/2 lung adenocarcinomas as two bands of 3.6 kb and 1.6 kb. No expression of L839P was observed in normal lung or trachea. No expression of L840P was observed in normal bone marrow, resting or activated PBMC, esophagus, or normal lung. Given the similar expression patterns, L839P and L840P may be derived from the same gene.

Further studies on L773P (SEQ ID NO: 58) resulted in the isolation of the extended consensus cDNA sequence provided in SEQ ID NO: 807.

Additional lung adenocarcinoma cDNA clones were isolated as follows. A cDNA library was prepared from a pool of two lung adenocarcinomas and subtracted
 5 against cDNA from a panel of normal tissues including lung, brain, liver, kidney, pancreas, skin, heart and spleen. The subtraction was performed using a PCR-based protocol (Clontech), which was modified to generate larger fragments. Within this protocol, tester and driver double stranded cDNA were separately digested with five restriction enzymes that recognize six-nucleotide restriction sites (MluI, MscI, PvuII, SalI and StuI). This
 10 digestion resulted in an average cDNA size of 600 bp, rather than the average size of 300 bp that results from digestion with RsaI according to the Clontech protocol. The ends of the restriction digested tester cDNA were filled in to generate blunt ends for adapter ligation. This modification did not affect the subtraction efficiency. Two tester populations were then created with different adapters, and the driver library remained
 15 without adapters. The tester and driver libraries were then hybridized using excess driver cDNA. In the first hybridization step, driver was separately hybridized with each of the two tester cDNA populations. This resulted in populations of (a) unhybridized tester cDNAs, (b) tester cDNAs hybridized to other tester cDNAs, (c) tester cDNAs hybridized to driver cDNAs and (d) unhybridized driver cDNAs. The two separate hybridization
 20 reactions were then combined, and rehybridized in the presence of additional denatured driver cDNA. Following this second hybridization, in addition to populations (a) through (d), a fifth population (e) was generated in which tester cDNA with one adapter hybridized to tester cDNA with the second adapter. Accordingly, the second hybridization step resulted in enrichment of differentially expressed sequences which could be used as
 25 templates for PCR amplification with adaptor-specific primers.

The ends were then filled in, and PCR amplification was performed using adaptor-specific primers. Only population (e), which contained tester cDNA that did not hybridize to driver cDNA, was amplified exponentially. A second PCR amplification step

was then performed, to reduce background and further enrich differentially expressed sequences.

Fifty-seven cDNA clones were isolated from the subtracted library (referred to as LAP1) and sequenced. The determined cDNA sequences for 16 of these clones are provided in SEQ ID NO: 101-116. The sequences of SEQ ID NO: 101 and 114 showed no significant homologies to previously identified sequences. The sequences of SEQ ID NO: 102-109 and 112 showed some similarity to previously identified sequences, while the sequences of SEQ ID NO: 113, 115 and 116 showed some similarity to previously isolated ESTs.

An additional 502 clones analyzed from the LAP1 library were sequenced and the determined cDNA sequences are shown in SEQ ID NO:828-1239 and 1564-1653.

C. Isolation of cDNA Sequences from Small Cell Lung Carcinoma Libraries using PCR-Based cDNA Library Subtraction

A subtracted cDNA library for small cell lung carcinoma (referred to as SCL1) was prepared using essentially the modified PCR-based subtraction process described above. cDNA from small cell lung carcinoma was subtracted against cDNA from a panel of normal tissues, including normal lung, brain, kidney, liver, pancreas, skin, heart, lymph node and spleen. Both tester and driver poly A⁺ RNA were initially amplified using SMART PCR cDNA synthesis kit (Clontech, Palo Alto, CA). The tester and driver double stranded cDNA were separately digested with five restriction enzymes (DraI, MscI, PvuII, SmaI, and StuI). These restriction enzymes generated blunt end cuts and the digestion resulted in an average insert size of 600 bp. Digestion with this set of restriction enzymes eliminates the step required to generate blunt ends by filling in of the cDNA ends. These modifications did not affect subtraction efficiency.

Eighty-five clones were isolated and sequenced. The determined cDNA sequences for 31 of these clones are provided in SEQ ID NO: 117-147. The sequences of SEQ ID NO: 122, 124, 126, 127, 130, 131, 133, 136, 139 and 147 showed no significant homologies to previously identified sequences. The sequences of SEQ ID NO: 120, 129,

135, 137, 140, 142, 144 and 145 showed some similarity to previously identified gene sequences, while the sequences of SEQ ID NO: 114, 118, 119, 121, 123, 125, 128, 132, 134, 138, 141, 143 and 147 showed some similarity to previously isolated ESTs.

In further studies, three additional cDNA libraries were generated from poly A+ RNA from a single small cell lung carcinoma sample subtracted against a pool of poly A+ RNA from nine normal tissues (lung, brain, kidney, liver, pancreas, skin, heart pituitary gland and spleen). For the first library (referred to as SCL2), the subtraction was carried out essentially as described above for the LAP1 library, with the exception that the tester and driver were digested with PvuII, StuI, MscI and DraI. The ratio of tester and driver cDNA used was as recommended by Clontech. For the second library (referred to as SCL3), subtraction was performed essentially as for SCL2 except that cDNA for highly redundant clones identified from the SCL2 library was included in the driver cDNA. Construction of the SCL4 library was performed essentially as described for the SCL3 library except that a higher ratio of driver to tester was employed.

Each library was characterized by DNA sequencing and database analyses. The determined cDNA sequence for 35 clones isolated from the SCL2 library are provided in SEQ ID NO: 245-279, with the determined cDNA sequences for 21 clones isolated from the SCL3 library and for 15 clones isolated from the SCL4 library being provided in SEQ ID NO: 280-300 and 301-315, respectively. The sequences of SEQ ID NO: 246, 254, 261, 262, 304, 309 and 311 showed no significant homologies to previously identified sequences. The sequence of SEQ ID NO: 245, 248, 255, 266, 270, 275, 280, 282, 283, 288-290, 292, 295, 301 and 303 showed some homology to previously isolated ESTs, while the sequences of SEQ ID NO: 247, 249-253, 256-260, 263-265, 267-269, 271-274, 276-279, 281, 284-287, 291, 293, 294, 296-300, 302, 305-308, 310 and 312-315 showed some homology to previously identified gene sequences.

3264 cDNA clones from three PCR-based subtracted cDNA libraries were analyzed by cDNA microarray technology as part of Lung Chip 5. Of the 3264 cDNA clones 960 clones came from SQL1 library, 768 clones came from SCL1 library, and 1536 clones came from SCL3 and SCL4 libraries. 35 pairs of fluorescent labeled cDNA probes

were used for the microarray analysis. Each probe pair included a lung tumor probe paired with a normal tissue probe. The expression data was analyzed. 498 cDNA clones were found to be overexpressed by 2-fold or greater in the small cell and/or non-small cell lung tumor probe groups compared to the normal tissue probe group. Also, the mean expression values for these clones in normal tissues were below 0.1 (range of expression is from 0.001 to 10). The cDNA sequences disclosed in SEQ ID NO:1240-1563 represent 324 non-redundant clones.

The following sequences were novel based on database analysis including GenBank and GeneSeq: SEQ ID NO:1240, 1243, 1247, 1269, 1272, 1280, 1283, 1285, 1286, 1289, 1300, 1309, 1318, 1319, 1327, 1335, 1339, 1346, 1359, 1369, 1370, 1371, 1393, 1398, 1405, 1408, 1413, 1414, 1417, 1422, 1429, 1432, 1435, 1436, 1438-1442, 1447, 1450, 1453, 1463, 1467, 1470, 1473, 1475, 1482, 1486, 1491-1494, 1501, 1505, 1506, 1514-1517, 1520, 1522, 1524, 1535, 1538, 1542, 1543, 1547, 1554, 1557, 1559, 1561, and 1563.

Full-length sequence for contig 139 (SEQ ID NO: 1467), also known as L985P, was identified by searching public databases using SEQ ID NO: 1467 as a query. By this approach, L985 was identified as cell surface immunomodulator-2 (CSIMM-2), the cDNA sequence of which is set forth in SEQ ID NO: 1676, encoding a protein having the sequence set forth in SEQ ID NO: 1677.

By microarray analysis, L985P was overexpressed by 2-fold or greater in the lung tumor probe groups compared to the normal tissue probe group. Moreover, the mean expression values for L985P in normal tissues was below 0.2 (range of expression was from 0.01 to 10). By real-time PCR analysis, greater than 40% of small cell lung carcinoma lung tumor samples tested had elevated L985P expression relative to normal lung tissue. Of those that displayed elevated L985P, the level of expression was greater than 3-fold higher than in normal lung tissue samples. Low or no expression of L985P was detected in an extensive panel of normal tissue RNAs. These findings for L985P support its use both as a diagnostic marker for detecting the presence of lung cancer in a patient and/or as an immunotherapeutic target for the treatment of lung cancer.

D. Isolation of cDNA Sequences from a Neuroendocrine Library using PCR-Based cDNA Library Subtraction

Using the modified PCR-based subtraction process, essentially as described above for the LAP1 subtracted library, a subtracted cDNA library (referred to as MLN1) was derived from a lung neuroendocrine carcinoma that had metastasized to the subcarinal lymph node, by subtraction with a panel of nine normal tissues, including normal lung, brain, kidney, liver, pancreas, skin, heart, lymph node and spleen.

Ninety-one individual clones were isolated and sequenced. The determined cDNA sequences for 58 of these clones are provided in SEQ ID NO: 147-222. The sequences of SEQ ID NO: 150, 151, 154, 157, 158, 159, 160, 163, 174, 175, 178, 186-190, 192, 193, 195-200, 208-210, 212-215 and 220 showed no significant homologies to previously identified sequences. The sequences of SEQ ID NO: 152, 155, 156, 161, 165, 166, 176, 179, 182, 184, 185, 191, 194, 221 and 222 showed some similarity to previously identified gene sequences, while the sequences of SEQ ID NO: 148, 149, 153, 164, 167-173, 177, 180, 181, 183, 201-207, 211 and 216-219 showed some similarity to previously isolated ESTs.

The determined cDNA sequences of an additional 442 clones isolated from the MLN1 library are provided in SEQ ID NO: 341-782. The determined cDNA sequences of an additional 11 clones isolated from the MLN1 library are provided in SEQ ID NO:1654-1664.

E. Isolation of cDNA Sequences from a Squamous Cell Lung Carcinoma Library using PCR-Based cDNA Library Subtraction

A subtracted cDNA library for squamous cell lung carcinoma (referred to as SQL1) was prepared, essentially using the modified PCR-based subtraction process described above, except the tester and driver double stranded cDNA were separately digested with four restriction enzymes (DraI, MscI, PvuII and StuI) cDNA from a pool of two squamous cell lung carcinomas was subtracted against cDNA from a pool of 10 normal

tissues, including normal lung, brain, kidney, liver, pancreas, skin, heart, spleen, esophagus and trachea.

Seventy-four clones were isolated and sequenced. The determined cDNA sequences for 22 of these clones are provided in SEQ ID NO: 223-244. The sequence of
 5 SEQ ID NO: 241 showed no significant homologies to previously identified sequences. The sequences of SEQ ID NO: 223, 225, 232, 233, 235, 238, 239, 242 and 243 showed some similarity to previously identified gene sequences, while the sequences of SEQ ID NO: 224, 226-231, 234, 236, 237, 240, 241 and 244 showed some similarity to previously isolated ESTs.

10 The sequences of an additional 12 clones isolated during characterization of cDNA libraries prepared from lung tumor tissue are provided in SEQ ID NO: 813-824. Comparison of these sequences with those in the GenBank database and the GeneSeq DNA database revealed no significant homologies to previously identified sequences.

15

EXAMPLE 2

SYNTHESIS OF POLYPEPTIDES

Polypeptides may be synthesized on a Perkin Elmer/Applied Biosystems Division 430A peptide synthesizer using FMOC chemistry with HPTU (O-Benzotriazole-
 20 N,N,N',N'-tetramethyluronium hexafluorophosphate) activation. A Gly-Cys-Gly sequence may be attached to the amino terminus of the peptide to provide a method of conjugation, binding to an immobilized surface, or labeling of the peptide. Cleavage of the peptides from the solid support may be carried out using the following cleavage mixture: trifluoroacetic acid:ethanedithiol:thioanisole:water :phenol (40:1:2:2:3). After cleaving for
 25 2 hours, the peptides may be precipitated in cold methyl-t-butyl-ether. The peptide pellets may then be dissolved in water containing 0.1% trifluoroacetic acid (TFA) and lyophilized prior to purification by C18 reverse phase HPLC. A gradient of 0%-60% acetonitrile (containing 0.1% TFA) in water (containing 0.1% TFA) may be used to elute the peptides.

Following lyophilization of the pure fractions, the peptides may be characterized using electrospray or other types of mass spectrometry and by amino acid analysis.

EXAMPLE 3

5 PREPARATION OF ANTIBODIES AGAINST LUNG CANCER ANTIGENS

Polyclonal antibodies against the lung cancer antigen L773P (SEQ ID NO: 783) were prepared as follows.

Rabbits were immunized with recombinant protein expressed in and purified
10 from *E. coli* as described above. For the initial immunization, 400 µg of antigen combined with muramyl dipeptide (MDP) was injected subcutaneously (S.C.). Animals were boosted S.C. 4 weeks later with 200 µg of antigen mixed with incomplete Freund's Adjuvant (IFA). Subsequent boosts of 100 µg of antigen mixed with IFA were injected S.C. as necessary to induce high antibody titer responses. Serum bleeds from immunized rabbits were tested for
15 L773P-specific reactivity using ELISA assays with purified protein and showed strong reactivity to L773P. Polyclonal antibodies against L773P were affinity purified from high titer polyclonal sera using purified protein attached to a solid support.

EXAMPLE 4

20 PROTEIN EXPRESSION OF LUNG TUMOR-SPECIFIC ANTIGENS

Full-length L773P (amino acids 2-364 of SEQ ID NO: 783), with a 6X His Tag, were subcloned into the pPDM expression vector and transformed into either BL21 CodonPlus or BL21 pLysS host cells using standard techniques. High levels of expression
25 were observed in both cases. Similarly, the N-terminal portion of L773P (amino acids 2-71 of SEQ ID NO: 783; referred to as L773PA), with a 6X His tag were subcloned into the vector pPDM and transformed into BL21 CodonPlus host cells. Low levels of expression were observed by N-terminal sequencing. The sequence of the expressed constructs for L773P and L773PA are provided in SEQ ID NO: 784 and 785, respectively.

EXAMPLE 5

EXPRESSION IN E. COLI OF L548S HIS TAG FUSION PROTEIN

5 The L548S coding region was PCR amplified with the following primers:

Forward primer starting at amino acid 2:

PDM-433: 5' gctaaaggtgaccccaagaaccaaag 3' Tm 60°C (SEQ ID NO:1665)

Reverse primer creating a XhoI site after the stop codon:

10 PDM-438: 5' ctattaactcgagggagacagataaacagtttcttta 3' Tm 61°C (SEQ ID NO:1666)

 The PCR product was then digested with XhoI restriction enzyme, gel purified and then cloned into pPDM His, a modified pET28 vector with a His tag in frame, which had been digested with Eco72I and XhoI restriction enzymes. The correct construct
15 was confirmed by DNA sequence analysis and then transformed into BL21 (DE3) pLys S and BL21 (DE3) CodonPlus RIL expression hosts.

 The protein sequence of expressed recombinant L548S is shown in SEQ ID NO:1667, and the DNA sequence of expressed recombinant L7548S is shown in
20 SEQ ID NO:1668.

 From the foregoing it will be appreciated that, although specific embodiments of the invention have been described herein for purposes of illustration, various modifications may be made without deviating from the spirit and scope of the
25 invention. Accordingly, the invention is not limited except as by the appended claims.

CLAIMS

What is claimed:

1. An isolated polypeptide, comprising at least an immunogenic portion of a lung tumor protein, wherein the tumor protein comprises an amino acid sequence that is encoded by a polynucleotide sequence selected from the group consisting of:

(a) sequences recited in SEQ ID NOs: 1, 11-13, 15, 20, 23-27, 29, 30, 33, 34, 39, 41, 43-46, 51, 52, 57, 58, 60, 62, 65-67, 69-71, 74, 76, 79, 80, 84, 86, 89-92, 95, 97, 98, 101, 110, 111, 113-119, 121-128, 130-134, 136, 138, 139, 141, 143, 146-151, 153, 154, 157-160, 162-164, 167-178, 180, 181, 183, 186-190, 192, 193, 195-220, 224, 226-231, 234, 236, 237, 240, 241, 244-246, 248, 254, 255, 261, 262, 266, 270, 275, 280, 282, 283, 288, 289, 290, 292, 295, 301, 303, 304, 309, 311, 341-782, 784, 785, 790, 792, 794, 796, 800, 802, 804, 807, 808, 811-826, 1240, 1243, 1247, 1269, 1272, 1280, 1283, 1285, 1286, 1289, 1300, 1309, 1318, 1319, 1327, 1335, 1339, 1346, 1359, 1369, 1370, 1371, 1393, 1398, 1405, 1408, 1413, 1414, 1417, 1422, 1429, 1432, 1435, 1436, 1438-1442, 1447, 1450, 1453, 1463, 1467, 1470, 1473, 1475, 1482, 1486, 1491-1494, 1501, 1505, 1506, 1514-1517, 1520, 1522, 1524, 1535, 1538, 1542, 1543, 1547, 1554, 1557, 1559, 1561, 1563 and 1669;

(b) sequences that hybridize to a sequence recited in any one of SEQ ID NOs: 1, 11-13, 15, 20, 23-27, 29, 30, 33, 34, 39, 41, 43-46, 51, 52, 57, 58, 60, 62, 65-67, 69-71, 74, 76, 79, 80, 84, 86, 89-92, 95, 97, 98, 101, 110, 111, 113-119, 121-128, 130-134, 136, 138, 139, 141, 143, 146-151, 153, 154, 157-160, 162-164, 167-178, 180, 181, 183, 186-190, 192, 193, 195-220, 224, 226-231, 234, 236, 237, 240, 241, 244-246, 248, 254, 255, 261, 262, 266, 270, 275, 280, 282, 283, 288, 289, 290, 292, 295, 301, 303, 304, 309, 311, 341-782, 784, 785, 790, 792, 794, 796, 800, 802, 804, 807, 808, 811-826, 1240, 1243, 1247, 1269, 1272, 1280, 1283, 1285, 1286, 1289, 1300, 1309, 1318, 1319, 1327, 1335, 1339, 1346, 1359, 1369, 1370, 1371, 1393, 1398, 1405, 1408, 1413, 1414, 1417, 1422, 1429, 1432, 1435, 1436, 1438-1442, 1447, 1450, 1453, 1463, 1467, 1470, 1473, 1475, 1482, 1486, 1491-1494, 1501, 1505, 1506, 1514-1517, 1520, 1522, 1524, 1535, 1538, 1542, 1543, 1547, 1554, 1557, 1559, 1561, 1563 and 1669 under moderately stringent conditions; and

(c) complements of sequences of (a) or (b).

2. An isolated polypeptide according to claim 1, wherein the polypeptide comprises an amino acid sequence that is encoded by a polynucleotide sequence recited in any one of SEQ ID NOs: 1, 11-13, 15, 20, 23-27, 29, 30, 33, 34, 39, 41, 43-46, 51, 52, 57, 58, 60, 62, 65-67, 69-71, 74, 76, 79, 80, 84, 86, 89-92, 95, 97, 98, 101, 110, 111, 113-119, 121-128, 130-134, 136, 138, 139, 141, 143, 146-151, 153, 154, 157-160, 162-164, 167-178, 180, 181, 183, 186-190, 192, 193, 195-220, 224, 226-231, 234, 236, 237, 240, 241, 244-246, 248, 254, 255, 261, 262, 266, 270, 275, 280, 282, 283, 288, 289, 290, 292, 295, 301, 303, 304, 309, 311, 341-782, 784, 785, 790, 792, 794, 796, 800, 802, 804, 807, 808, 811-826, 1240, 1243, 1247, 1269, 1272, 1280, 1283, 1285, 1286, 1289, 1300, 1309, 1318, 1319, 1327, 1335, 1339, 1346, 1359, 1369, 1370, 1371, 1393, 1398, 1405, 1408, 1413, 1414, 1417, 1422, 1429, 1432, 1435, 1436, 1438-1442, 1447, 1450, 1453, 1463, 1467, 1470, 1473, 1475, 1482, 1486, 1491-1494, 1501, 1505, 1506, 1514-1517, 1520, 1522, 1524, 1535, 1538, 1542, 1543, 1547, 1554, 1557, 1559, 1561, 1563, and 1669 or a complement of any of the foregoing polynucleotide sequences.

3. An isolated polypeptide comprising a sequence recited in any one of SEQ ID NOs: 786, 787, 791, 793, 795, 797-799, 806, 809, 827 and 1670-1675.

4. An isolated polynucleotide encoding at least 15 amino acid residues of a lung tumor protein, or a variant thereof that differs in one or more substitutions, deletions, additions and/or insertions such that the ability of the variant to react with antigen-specific antisera is not substantially diminished, wherein the tumor protein comprises an amino acid sequence that is encoded by a polynucleotide comprising a sequence recited in any one of SEQ ID NO: 1, 11-13, 15, 20, 23-27, 29, 30, 33, 34, 39, 41, 43-46, 51, 52, 57, 58, 60, 62, 65-67, 69-71, 74, 76, 79, 80, 84, 86, 89-92, 95, 97, 98, 101, 110, 111, 113-119, 121-128, 130-134, 136, 138, 139, 141, 143, 146-151, 153, 154, 157-160, 162-164, 167-178, 180, 181, 183, 186-190, 192, 193, 195-220, 224, 226-231, 234, 236, 237, 240, 241, 244-246, 248, 254, 255, 261, 262, 266, 270, 275, 280, 282, 283, 288, 289, 290, 292, 295, 301, 303, 304, 309, 311, 341-782, 784, 785,

790, 792, 794, 796, 800, 802, 804, 807, 808, 811-826, 1240, 1243, 1247, 1269, 1272, 1280, 1283, 1285, 1286, 1289, 1300, 1309, 1318, 1319, 1327, 1335, 1339, 1346, 1359, 1369, 1370, 1371, 1393, 1398, 1405, 1408, 1413, 1414, 1417, 1422, 1429, 1432, 1435, 1436, 1438-1442, 1447, 1450, 1453, 1463, 1467, 1470, 1473, 1475, 1482, 1486, 1491-1494, 1501, 1505, 1506, 1514-1517, 1520, 1522, 1524, 1535, 1538, 1542, 1543, 1547, 1554, 1557, 1559, 1561, 1563 and 1669 or a complement of any of the foregoing sequences.

5. An isolated polynucleotide encoding a lung tumor protein, or a variant thereof, wherein the tumor protein comprises an amino acid sequence that is encoded by a polynucleotide comprising a sequence recited in any one of SEQ ID NOs: 1, 11-13, 15, 20, 23-27, 29, 30, 33, 34, 39, 41, 43-46, 51, 52, 57, 58, 60, 62, 65-67, 69-71, 74, 76, 79, 80, 84, 86, 89-92, 95, 97, 98, 101, 110, 111, 113-119, 121-128, 130-134, 136, 138, 139, 141, 143, 146-151, 153, 154, 157-160, 162-164, 167-178, 180, 181, 183, 186-190, 192, 193, 195-220, 224, 226-231, 234, 236, 237, 240, 241, 244-246, 248, 254, 255, 261, 262, 266, 270, 275, 280, 282, 283, 288, 289, 290, 292, 295, 301, 303, 304, 309, 311, 341-782, 784, 785, 790, 792, 794, 796, 800, 802, 804, 807, 808, 811-826, 1240, 1243, 1247, 1269, 1272, 1280, 1283, 1285, 1286, 1289, 1300, 1309, 1318, 1319, 1327, 1335, 1339, 1346, 1359, 1369, 1370, 1371, 1393, 1398, 1405, 1408, 1413, 1414, 1417, 1422, 1429, 1432, 1435, 1436, 1438-1442, 1447, 1450, 1453, 1463, 1467, 1470, 1473, 1475, 1482, 1486, 1491-1494, 1501, 1505, 1506, 1514-1517, 1520, 1522, 1524, 1535, 1538, 1542, 1543, 1547, 1554, 1557, 1559, 1561, 1563 and 1669 or a complement of any of the foregoing sequences.

6. An isolated polynucleotide, comprising a sequence recited in any one of SEQ ID NOs: 1, 11-13, 15, 20, 23-27, 29, 30, 33, 34, 39, 41, 43-46, 51, 52, 57, 58, 60, 62, 65-67, 69-71, 74, 76, 79, 80, 84, 86, 89-92, 95, 97, 98, 101, 110, 111, 113-119, 121-128, 130-134, 136, 138, 139, 141, 143, 146-151, 153, 154, 157-160, 162-164, 167-178, 180, 181, 183, 186-190, 192, 193, 195-220, 224, 226-231, 234, 236, 237, 240, 241, 244-246, 248, 254, 255, 261, 262, 266, 270, 275, 280, 282, 283, 288, 289, 290, 292, 295, 301, 303, 304, 309, 311, 341-782, 784, 785, 790, 792, 794, 796, 800, 802, 804, 807, 808, 811-826, 1240, 1243, 1247, 1269, 1272, 1280,

1283, 1285, 1286, 1289, 1300, 1309, 1318, 1319, 1327, 1335, 1339, 1346, 1359, 1369, 1370, 1371, 1393, 1398, 1405, 1408, 1413, 1414, 1417, 1422, 1429, 1432, 1435, 1436, 1438-1442, 1447, 1450, 1453, 1463, 1467, 1470, 1473, 1475, 1482, 1486, 1491-1494, 1501, 1505, 1506, 1514-1517, 1520, 1522, 1524, 1535, 1538, 1542, 1543, 1547, 1554, 1557, 1559, 1561, 1563 and 1669.

7. An isolated polynucleotide, comprising a sequence that hybridizes to a sequence recited in any one of SEQ ID NOs: 1, 11-13, 15, 20, 23-27, 29, 30, 33, 34, 39, 41, 43-46, 51, 52, 57, 58, 60, 62, 65-67, 69-71, 74, 76, 79, 80, 84, 86, 89-92, 95, 97, 98, 101, 110, 111, 113-119, 121-128, 130-134, 136, 138, 139, 141, 143, 146-151, 153, 154, 157-160, 162-164, 167-178, 180, 181, 183, 186-190, 192, 193, 195-220, 224, 226-231, 234, 236, 237, 240, 241, 244-246, 248, 254, 255, 261, 262, 266, 270, 275, 280, 282, 283, 288, 289, 290, 292, 295, 301, 303, 304, 309, 311, 341-782, 784, 785, 790, 792, 794, 796, 800, 802, 804, 807, 808, 811-826, 1240, 1243, 1247, 1269, 1272, 1280, 1283, 1285, 1286, 1289, 1300, 1309, 1318, 1319, 1327, 1335, 1339, 1346, 1359, 1369, 1370, 1371, 1393, 1398, 1405, 1408, 1413, 1414, 1417, 1422, 1429, 1432, 1435, 1436, 1438-1442, 1447, 1450, 1453, 1463, 1467, 1470, 1473, 1475, 1482, 1486, 1491-1494, 1501, 1505, 1506, 1514-1517, 1520, 1522, 1524, 1535, 1538, 1542, 1543, 1547, 1554, 1557, 1559, 1561, 1563 and 1669 under moderately stringent conditions.

8. An isolated polynucleotide complementary to a polynucleotide according to any one of claims 4-7.

9. An expression vector, comprising a polynucleotide according to any one of claims 4-8.

10. A host cell transformed or transfected with an expression vector according to claim 9.

11. An isolated antibody, or antigen-binding fragment thereof, that specifically binds to a lung tumor protein that comprises an amino acid sequence that is encoded by a polynucleotide sequence recited in any one of SEQ ID NOs: 1, 11-13, 15, 20, 23-27, 29, 30, 33, 34, 39, 41, 43-46, 51, 52, 57, 58, 60, 62, 65-67, 69-71, 74, 76, 79, 80, 84, 86, 89-92, 95, 97, 98, 101, 110, 111, 113-119, 121-128, 130-134, 136, 138, 139, 141, 143, 146-151, 153, 154, 157-160, 162-164, 167-178, 180, 181, 183, 186-190, 192, 193, 195-220, 224, 226-231, 234, 236, 237, 240, 241, 244-246, 248, 254, 255, 261, 262, 266, 270, 275, 280, 282, 283, 288, 289, 290, 292, 295, 301, 303, 304, 309, 311, 341-782, 784, 785, 790, 792, 794, 796, 800, 802, 804, 807, 808, 811-826, 1240, 1243, 1247, 1269, 1272, 1280, 1283, 1285, 1286, 1289, 1300, 1309, 1318, 1319, 1327, 1335, 1339, 1346, 1359, 1369, 1370, 1371, 1393, 1398, 1405, 1408, 1413, 1414, 1417, 1422, 1429, 1432, 1435, 1436, 1438-1442, 1447, 1450, 1453, 1463, 1467, 1470, 1473, 1475, 1482, 1486, 1491-1494, 1501, 1505, 1506, 1514-1517, 1520, 1522, 1524, 1535, 1538, 1542, 1543, 1547, 1554, 1557, 1559, 1561, 1563 and 1669 or a complement of any of the foregoing polynucleotide sequences.

12. A fusion protein, comprising at least one polypeptide according to claim 1.

13. A fusion protein according to claim 12, wherein the fusion protein comprises an expression enhancer that increases expression of the fusion protein in a host cell transfected with a polynucleotide encoding the fusion protein.

14. A fusion protein according to claim 12, wherein the fusion protein comprises a T helper epitope that is not present within the polypeptide of claim 1.

15. A fusion protein according to claim 12, wherein the fusion protein comprises an affinity tag.

16. An isolated polynucleotide encoding a fusion protein according to claim 12.

17. A pharmaceutical composition, comprising a physiologically acceptable carrier and at least one component selected from the group consisting of:

- (a) a polypeptide according to claim 1;
- (b) a polynucleotide according to claim 4;
- (c) an antibody according to claim 11;
- (d) a fusion protein according to claim 12; and
- (e) a polynucleotide according to claim 16.

18. An immunogenic composition comprising an immunostimulant and at least one component selected from the group consisting of:

- (a) a polypeptide according to claim 1;
- (b) a polynucleotide according to claim 4;
- (c) an antibody according to claim 11;
- (d) a fusion protein according to claim 12; and
- (e) a polynucleotide according to claim 16.

19. An immunogenic composition according to claim 18, wherein the immunostimulant is an adjuvant.

20. An immunogenic composition according to any claim 18, wherein the immunostimulant induces a predominantly Type I response.

21. A method for inhibiting the development of a cancer in a patient, comprising administering to a patient an effective amount of a pharmaceutical composition according to claim 17.

22. A method for inhibiting the development of a cancer in a patient, comprising administering to a patient an effective amount of an immunogenic composition according to claim 18.

23. A pharmaceutical composition comprising an antigen-presenting cell that expresses a polypeptide according to claim 1, in combination with a pharmaceutically acceptable carrier or excipient.

24. A pharmaceutical composition according to claim 23, wherein the antigen presenting cell is a dendritic cell or a macrophage.

25. An immunogenic composition comprising an antigen-presenting cell that expresses a polypeptide comprising at least an immunogenic portion of a lung tumor protein, or a variant thereof, wherein the tumor protein comprises an amino acid sequence that is encoded by a polynucleotide sequence selected from the group consisting of:

(a) sequences recited in SEQ ID NOs: 1, 11-13, 15, 20, 23-27, 29, 30, 33, 34, 39, 41, 43-46, 51, 52, 57, 58, 60, 62, 65-67, 69-71, 74, 76, 79, 80, 84, 86, 89-92, 95, 97, 98, 101, 110, 111, 113-119, 121-128, 130-134, 136, 138, 139, 141, 143, 146-151, 153, 154, 157-160, 162-164, 167-178, 180, 181, 183, 186-190, 192, 193, 195-220, 224, 226-231, 234, 236, 237, 240, 241, 244-246, 248, 254, 255, 261, 262, 266, 270, 275, 280, 282, 283, 288, 289, 290, 292, 295, 301, 303, 304, 309, 311, 341-782, 784, 785, 790, 792, 794, 796, 800-804, 807, 808, 810-826 and 828-1664, 1669 and 1676;

(b) sequences that hybridize to a sequence recited in any one of SEQ ID NOs: 1, 11-13, 15, 20, 23-27, 29, 30, 33, 34, 39, 41, 43-46, 51, 52, 57, 58, 60, 62, 65-67, 69-71, 74, 76, 79, 80, 84, 86, 89-92, 95, 97, 98, 101, 110, 111, 113-119, 121-128, 130-134, 136, 138, 139, 141, 143, 146-151, 153, 154, 157-160, 162-164, 167-178, 180, 181, 183, 186-190, 192, 193, 195-220, 224, 226-231, 234, 236, 237, 240, 241, 244-246, 248, 254, 255, 261, 262, 266, 270, 275, 280, 282, 283, 288, 289, 290, 292, 295, 301, 303, 304, 309, 311, 341-782, 784, 785, 790, 792, 794, 796, 800-804, 807, 808, 810-826, 828-1664, 1669 and 1676 under moderately stringent conditions; and

(c) complements of sequences of (i) or (ii);

in combination with an immunostimulant.

26. An immunogenic composition according to claim 25, wherein the immunostimulant is an adjuvant.

27. An immunogenic composition according to claim 25, wherein the immunostimulant induces a predominantly Type I response.

28. An immunogenic composition according to claim 25, wherein the antigen-presenting cell is a dendritic cell.

29. A method for inhibiting the development of a cancer in a patient, comprising administering to a patient an effective amount of an antigen-presenting cell that expresses a polypeptide comprising at least an immunogenic portion of a lung tumor protein, or a variant thereof, wherein the tumor protein comprises an amino acid sequence that is encoded by a polynucleotide sequence selected from the group consisting of:

(a) sequences recited in SEQ ID NOs: 1, 11-13, 15, 20, 23-27, 29, 30, 33, 34, 39, 41, 43-46, 51, 52, 57, 58, 60, 62, 65-67, 69-71, 74, 76, 79, 80, 84, 86, 89-92, 95, 97, 98, 101, 110, 111, 113-119, 121-128, 130-134, 136, 138, 139, 141, 143, 146-151, 153, 154, 157-160, 162-164, 167-178, 180, 181, 183, 186-190, 192, 193, 195-220, 224, 226-231, 234, 236, 237, 240, 241, 244-246, 248, 254, 255, 261, 262, 266, 270, 275, 280, 282, 283, 288, 289, 290, 292, 295, 301, 303, 304, 309, 311, 341-782, 784, 785, 790, 792, 794, 796, 800-804, 807, 808, 810-826 828-1664, 1669 and 1676;

(b) sequences that hybridize to a sequence recited in any one of SEQ ID NOs: 1, 11-13, 15, 20, 23-27, 29, 30, 33, 34, 39, 41, 43-46, 51, 52, 57, 58, 60, 62, 65-67, 69-71, 74, 76, 79, 80, 84, 86, 89-92, 95, 97, 98, 101, 110, 111, 113-119, 121-128, 130-134, 136, 138, 139, 141, 143, 146-151, 153, 154, 157-160, 162-164, 167-178, 180, 181, 183, 186-190, 192, 193, 195-220, 224, 226-231, 234, 236, 237, 240, 241, 244-246, 248, 254, 255, 261, 262, 266, 270, 275, 280, 282, 283, 288, 289, 290, 292, 295, 301, 303, 304, 309, 311, 341-782, 784, 785, 790, 792, 794, 796, 800-804, 807, 808, 810-826 828-1664, 1669 and 1676 under moderately stringent conditions; and

(c) complements of sequences of (i) or (ii) encoded by a polynucleotide recited in any one of SEQ ID NOs: 1, 11-13, 15, 20, 23-27, 29, 30, 33, 34, 39, 41, 43-46, 51, 52, 57, 58, 60, 62, 65-67, 69-71, 74, 76, 79, 80, 84, 86, 89-92, 95, 97, 98, 101, 110, 111, 113-119, 121-128, 130-134, 136, 138, 139, 141, 143, 146-151, 153, 154, 157-160, 162-164, 167-178, 180, 181, 183, 186-190, 192, 193, 195-220, 224, 226-231, 234, 236, 237, 240, 241, 244-246, 248, 254, 255, 261, 262, 266, 270, 275, 280, 282, 283, 288, 289, 290, 292, 295, 301, 303, 304, 309, 311, 341-782, 784, 785, 790, 792, 794, 796, 800-804, 807, 808, 810-826 828-1664, 1669 and 1676;

and thereby inhibiting the development of a cancer in the patient.

30. A method according to claim 29, wherein the antigen-presenting cell is a dendritic cell.

31. A method according to any one of claims 21, 22 and 29, wherein the cancer is lung cancer.

32. A method for removing tumor cells from a biological sample, comprising contacting a biological sample with T cells that specifically react with a lung tumor protein, wherein the tumor protein comprises an amino acid sequence that is encoded by a polynucleotide sequence selected from the group consisting of:

(i) polynucleotides recited in any one of SEQ ID NOs: 1, 11-13, 15, 20, 23-27, 29, 30, 33, 34, 39, 41, 43-46, 51, 52, 57, 58, 60, 62, 65-67, 69-71, 74, 76, 79, 80, 84, 86, 89-92, 95, 97, 98, 101, 110, 111, 113-119, 121-128, 130-134, 136, 138, 139, 141, 143, 146-151, 153, 154, 157-160, 162-164, 167-178, 180, 181, 183, 186-190, 192, 193, 195-220, 224, 226-231, 234, 236, 237, 240, 241, 244-246, 248, 254, 255, 261, 262, 266, 270, 275, 280, 282, 283, 288, 289, 290, 292, 295, 301, 303, 304, 309, 311, 341-782, 784, 785, 790, 792, 794, 796, 800-804, 807, 808, 810-826 828-1664, 1669 and 1676 ; and

(ii) complements of the foregoing polynucleotides;

wherein the step of contacting is performed under conditions and for a time sufficient to permit the removal of cells expressing the antigen from the sample.

33. A method according to claim 32, wherein the biological sample is blood or a fraction thereof.

34. A method for inhibiting the development of a cancer in a patient, comprising administering to a patient a biological sample treated according to the method of claim 32.

35. A method for stimulating and/or expanding T cells specific for a lung tumor protein, comprising contacting T cells with at least one component selected from the group consisting of:

(a) polypeptides comprising at least an immunogenic portion of a lung tumor protein, or a variant thereof, wherein the tumor protein comprises an amino acid sequence that is encoded by a polynucleotide sequence selected from the group consisting of:

(i) sequences recited in SEQ ID NOs: 1, 11-13, 15, 20, 23-27, 29, 30, 33, 34, 39, 41, 43-46, 51, 52, 57, 58, 60, 62, 65-67, 69-71, 74, 76, 79, 80, 84, 86, 89-92, 95, 97, 98, 101, 110, 111, 113-119, 121-128, 130-134, 136, 138, 139, 141, 143, 146-151, 153, 154, 157-160, 162-164, 167-178, 180, 181, 183, 186-190, 192, 193, 195-220, 224, 226-231, 234, 236, 237, 240, 241, 244-246, 248, 254, 255, 261, 262, 266, 270, 275, 280, 282, 283, 288, 289, 290, 292, 295, 301, 303, 304, 309, 311, 341-782, 784, 785, 790, 792, 794, 796, 800-804, 807, 808, 810-826, 828-1664, 1669 and 1676;

(ii) sequences that hybridize to a sequence recited in any one of SEQ ID NOs: 1, 11-13, 15, 20, 23-27, 29, 30, 33, 34, 39, 41, 43-46, 51, 52, 57, 58, 60, 62, 65-67, 69-71, 74, 76, 79, 80, 84, 86, 89-92, 95, 97, 98, 101, 110, 111, 113-119, 121-128, 130-134, 136, 138, 139, 141, 143, 146-151, 153, 154, 157-160, 162-164, 167-178, 180, 181, 183, 186-190, 192, 193, 195-220, 224, 226-231, 234, 236, 237, 240, 241, 244-246, 248, 254, 255, 261, 262, 266, 270, 275, 280, 282, 283, 288, 289, 290, 292, 295, 301, 303, 304, 309, 311, 341-782, 784, 785,

790, 792, 794, 796, 800-804, 807, 808, 810-826 828-1664, 1669 and 1676 under moderately stringent conditions; and

- (iii) complements of sequences of (i) or (ii);
- (b) polynucleotides encoding a polypeptide of (a); and
- (c) antigen presenting cells that express a polypeptide of (a);

under conditions and for a time sufficient to permit the stimulation and/or expansion of T cells.

36. An isolated T cell population, comprising T cells prepared according to the method of claim 35.

37. A method for inhibiting the development of a cancer in a patient, comprising administering to a patient an effective amount of a T cell population according to claim 36.

38. A method for inhibiting the development of a cancer in a patient, comprising the steps of:

(a) incubating CD4⁺ and/or CD8⁺ T cells isolated from a patient with at least one component selected from the group consisting of:

(i) polypeptides comprising at least an immunogenic portion of a lung tumor protein, or a variant thereof, wherein the tumor protein comprises an amino acid sequence that is encoded by a polynucleotide sequence selected from the group consisting of:

(1) sequences recited in SEQ ID NOs: 1, 11-13, 15, 20, 23-27, 29, 30, 33, 34, 39, 41, 43-46, 51, 52, 57, 58, 60, 62, 65-67, 69-71, 74, 76, 79, 80, 84, 86, 89-92, 95, 97, 98, 101, 110, 111, 113-119, 121-128, 130-134, 136, 138, 139, 141, 143, 146-151, 153, 154, 157-160, 162-164, 167-178, 180, 181, 183, 186-190, 192, 193, 195-220, 224, 226-231, 234, 236, 237, 240, 241, 244-246, 248, 254, 255, 261, 262, 266, 270, 275, 280, 282, 283, 288, 289, 290, 292, 295, 301, 303, 304, 309, 311, 341-782, 784, 785, 790, 792, 794, 796, 800-804, 807, 808, 810-826, 828-1664, 1669 and 1676;

(3) complements of sequences of (1) or (2);

(ii) polynucleotides encoding a polypeptide of (i); and

(iii) antigen presenting cells that expresses a polypeptide of (i);

such that T cells proliferate; and

(b) administering to the patient an effective amount of the proliferated T cells,

and thereby inhibiting the development of a cancer in the patient.

(a) incubating CD4⁺ and/or CD8⁺ T cells isolated from a patient with at least one component selected from the group consisting of:

(1) sequences recited in SEQ ID NOs: 1, 11-13, 15, 20, 23-27, 29, 30, 33, 34, 39, 41, 43-46, 51, 52, 57, 58, 60, 62, 65-67, 69-71, 74, 76, 79, 80, 84, 86, 89-92, 95, 97, 98, 101, 110, 111, 113-119, 121-128, 130-134, 136, 138, 139, 141, 143, 146-151, 153, 154, 157-160, 162-164, 167-178, 180, 181, 183, 186-190, 192, 193, 195-220, 224, 226-231, 234, 236, 237, 240, 241, 244-246, 248, 254, 255, 261, 262, 266, 270, 275, 280, 282, 283, 288, 289, 290, 292, 295, 301, 303, 304, 309, 311, 341-782, 784, 785, 790, 792, 794, 796, 800-804, 807, 808, 810-826, 828-1664, 1669 and 1676;

(2) sequences that hybridize to a sequence recited in any one of SEQ ID NOs: 1, 11-13, 15, 20, 23-27, 29, 30, 33, 34, 39, 41, 43-46, 51, 52, 57, 58, 60, 62, 65-67, 69-71, 74, 76, 79, 80, 84, 86, 89-92, 95, 97, 98, 101, 110, 111, 113-119, 121-128, 130-134, 136, 138, 139, 141, 143, 146-151, 153, 154, 157-160, 162-164, 167-178, 180, 181, 183, 186-190, 192, 193, 195-220, 224, 226-231, 234, 236, 237, 240, 241, 244-246, 248, 254, 255, 261, 262, 266, 270, 275, 280, 282, 283, 288, 289, 290, 292, 295, 301, 303, 304, 309, 311, 341-782, 784, 785, 790, 792, 794, 796, 800-804, 807, 808, 810-826, 828-1664, 1669 and 1676 under moderately stringent conditions; and

(3) complements of sequences of (1) or (2);

(ii) polynucleotides encoding a polypeptide of (i); and

(iii) antigen presenting cells that express a polypeptide of (i);

such that T cells proliferate;

(b) cloning at least one proliferated cell to provide cloned T cells; and

(c) administering to the patient an effective amount of the cloned T cells, and thereby inhibiting the development of a cancer in the patient.

40. A method for determining the presence or absence of a cancer in a patient, comprising the steps of:

(a) contacting a biological sample obtained from a patient with a binding agent that binds to a lung tumor protein, wherein the tumor protein comprises an amino acid sequence that is encoded by a polynucleotide sequence recited in any one of SEQ ID NOs: 1, 11-13, 15, 20, 23-27, 29, 30, 33, 34, 39, 41, 43-46, 51, 52, 57, 58, 60, 62, 65-67, 69-71, 74, 76, 79, 80, 84, 86, 89-92, 95, 97, 98, 101, 110, 111, 113-119, 121-128, 130-134, 136, 138, 139, 141, 143, 146-151, 153, 154, 157-160, 162-164, 167-178, 180, 181, 183, 186-190, 192, 193, 195-220, 224, 226-231, 234, 236, 237, 240, 241, 244-246, 248, 254, 255, 261, 262, 266, 270, 275, 280, 282, 283, 288, 289, 290, 292, 295, 301, 303, 304, 309, 311, 341-782, 784, 785, 790, 792, 794, 796, 800-804, 807, 808, 810-826, 828-1664, 1669 and 1676 or a complement of any of the foregoing polynucleotide sequences;

(b) detecting in the sample an amount of polypeptide that binds to the binding agent; and

(c) comparing the amount of polypeptide to a predetermined cut-off value, and therefrom determining the presence or absence of a cancer in the patient.

41. A method according to claim 40, wherein the binding agent is an antibody.

42. A method according to claim 43, wherein the antibody is a monoclonal antibody.

43. A method according to claim 40, wherein the cancer is lung cancer.

44. A method for monitoring the progression of a cancer in a patient, comprising the steps of:

(a) contacting a biological sample obtained from a patient at a first point in time with a binding agent that binds to a lung tumor protein, wherein the tumor protein comprises an amino acid sequence that is encoded by a polynucleotide sequence recited in any one of SEQ ID NOs: 1, 11-13, 15, 20, 23-27, 29, 30, 33, 34, 39, 41, 43-46, 51, 52, 57, 58, 60, 62, 65-67, 69-71, 74, 76, 79, 80, 84, 86, 89-92, 95, 97, 98, 101, 110, 111, 113-119, 121-128, 130-134, 136, 138, 139, 141, 143, 146-151, 153, 154, 157-160, 162-164, 167-178, 180, 181, 183, 186-190, 192, 193, 195-220, 224, 226-231, 234, 236, 237, 240, 241, 244-246, 248, 254, 255, 261, 262, 266, 270, 275, 280, 282, 283, 288, 289, 290, 292, 295, 301, 303, 304, 309, 311, 341-782, 784, 785, 790, 792, 794, 796, 800-804, 807, 808, 810-826, 828-1664, 1669 and 1676 or a complement of any of the foregoing polynucleotide sequences;

(b) detecting in the sample an amount of polypeptide that binds to the binding agent;

(c) repeating steps (a) and (b) using a biological sample obtained from the patient at a subsequent point in time; and

(d) comparing the amount of polypeptide detected in step (c) to the amount detected in step (b) and therefrom monitoring the progression of the cancer in the patient.

45. A method according to claim 44, wherein the binding agent is an antibody.

46. A method according to claim 45, wherein the antibody is a monoclonal antibody.

47. A method according to claim 44, wherein the cancer is a lung cancer.

48. A method for determining the presence or absence of a cancer in a patient, comprising the steps of:

(a) contacting a biological sample obtained from a patient with an oligonucleotide that hybridizes to a polynucleotide that encodes a lung tumor protein, wherein the tumor protein comprises an amino acid sequence that is encoded by a polynucleotide sequence recited in any one of SEQ ID NO: 1, 11-13, 15, 20, 23-27, 29, 30, 33, 34, 39, 41, 43-46, 51, 52, 57, 58, 60, 62, 65-67, 69-71, 74, 76, 79, 80, 84, 86, 89-92, 95, 97, 98, 101, 110, 111, 113-119, 121-128, 130-134, 136, 138, 139, 141, 143, 146-151, 153, 154, 157-160, 162-164, 167-178, 180, 181, 183, 186-190, 192, 193, 195-220, 224, 226-231, 234, 236, 237, 240, 241, 244-246, 248, 254, 255, 261, 262, 266, 270, 275, 280, 282, 283, 288, 289, 290, 292, 295, 301, 303, 304, 309, 311, 341-782, 784, 785, 790, 792, 794, 796, 800-804, 807, 808, 810-826, 828-1664, 1669 and 1676 or a complement of any of the foregoing polynucleotide sequences;

(b) detecting in the sample an amount of a polynucleotide that hybridizes to the oligonucleotide; and

(c) comparing the amount of polynucleotide that hybridizes to the oligonucleotide to a predetermined cut-off value, and therefrom determining the presence or absence of a cancer in the patient.

49. A method according to claim 48, wherein the amount of polynucleotide that hybridizes to the oligonucleotide is determined using a polymerase chain reaction.

50. A method according to claim 48, wherein the amount of polynucleotide that hybridizes to the oligonucleotide is determined using a hybridization assay.

51. A method for monitoring the progression of a cancer in a patient, comprising the steps of:

(a) contacting a biological sample obtained from a patient with an oligonucleotide that hybridizes to a polynucleotide that encodes a lung tumor protein, wherein the tumor protein comprises an amino acid sequence that is encoded by a polynucleotide sequence recited in any one of SEQ ID NO: 1, 11-13, 15, 20, 23-27, 29, 30, 33, 34, 39, 41, 43-46, 51, 52, 57, 58, 60, 62, 65-67, 69-71, 74, 76, 79, 80, 84, 86, 89-92, 95, 97, 98, 101, 110, 111, 113-119, 121-128, 130-134, 136, 138, 139, 141, 143, 146-151, 153, 154, 157-160, 162-164, 167-178, 180, 181, 183, 186-190, 192, 193, 195-220, 224, 226-231, 234, 236, 237, 240, 241, 244-246, 248, 254, 255, 261, 262, 266, 270, 275, 280, 282, 283, 288, 289, 290, 292, 295, 301, 303, 304, 309, 311, 341-782, 784, 785, 790, 792, 794, 796, 800-804, 807, 808, 810-826, 828-1664, 1669 and 1676 or a complement of any of the foregoing polynucleotide sequences;

(b) detecting in the sample an amount of a polynucleotide that hybridizes to the oligonucleotide;

(c) repeating steps (a) and (b) using a biological sample obtained from the patient at a subsequent point in time; and

(d) comparing the amount of polynucleotide detected in step (c) to the amount detected in step (b) and therefrom monitoring the progression of the cancer in the patient.

52. A method according to claim 51, wherein the amount of polynucleotide that hybridizes to the oligonucleotide is determined using a polymerase chain reaction.

53. A method according to claim 51, wherein the amount of polynucleotide that hybridizes to the oligonucleotide is determined using a hybridization assay.

54. A diagnostic kit, comprising:

- (a) one or more antibodies according to claim 11; and
- (b) a detection reagent comprising a reporter group.

55. A kit according to claim 54, wherein the antibodies are immobilized on a solid support.

56. A kit according to claim 54, wherein the detection reagent comprises an anti-immunoglobulin, protein G, protein A or lectin.

57. A kit according to claim 54, wherein the reporter group is selected from the group consisting of radioisotopes, fluorescent groups, luminescent groups, enzymes, biotin and dye particles.

58. An oligonucleotide comprising 10 to 40 contiguous nucleotides that hybridize under moderately stringent conditions to a polynucleotide that encodes a lung tumor protein, wherein the tumor protein comprises an amino acid sequence that is encoded by a polynucleotide sequence recited in any one of SEQ ID NOs: 1, 11-13, 15, 20, 23-27, 29, 30, 33, 34, 39, 41, 43-46, 51, 52, 57, 58, 60, 62, 65-67, 69-71, 74, 76, 79, 80, 84, 86, 89-92, 95, 97, 98, 101, 110, 111, 113-119, 121-128, 130-134, 136, 138, 139, 141, 143, 146-151, 153, 154, 157-160, 162-164, 167-178, 180, 181, 183, 186-190, 192, 193, 195-220, 224, 226-231, 234, 236, 237, 240, 241, 244-246, 248, 254, 255, 261, 262, 266, 270, 275, 280, 282, 283, 288, 289, 290, 292, 295, 301, 303, 304, 309, 311, 341-782, 784, 785, 790, 792, 794, 796, 800, 802, 804, 807, 808, 811-826, 1240, 1243, 1247, 1269, 1272, 1280, 1283, 1285, 1286, 1289, 1300, 1309, 1318, 1319, 1327, 1335, 1339, 1346, 1359, 1369, 1370, 1371, 1393, 1398, 1405, 1408, 1413, 1414, 1417, 1422, 1429, 1432, 1435, 1436, 1438-1442, 1447, 1450, 1453, 1463, 1467, 1470, 1473, 1475,

1482, 1486, 1491-1494, 1501, 1505, 1506, 1514-1517, 1520, 1522, 1524, 1535, 1538, 1542, 1543, 1547, 1554, 1557, 1559, 1561, 1563, 1669 and 1676 or a complement of any of the foregoing polynucleotides.

59. A oligonucleotide according to claim 58, wherein the oligonucleotide comprises 10-40 contiguous nucleotides recited in any one of SEQ ID NO: 1, 11-13, 15, 20, 23-27, 29, 30, 33, 34, 39, 41, 43-46, 51, 52, 57, 58, 60, 62, 65-67, 69-71, 74, 76, 79, 80, 84, 86, 89-92, 95, 97, 98, 101, 110, 111, 113-119, 121-128, 130-134, 136, 138, 139, 141, 143, 146-151, 153, 154, 157-160, 162-164, 167-178, 180, 181, 183, 186-190, 192, 193, 195-220, 224, 226-231, 234, 236, 237, 240, 241, 244-246, 248, 254, 255, 261, 262, 266, 270, 275, 280, 282, 283, 288, 289, 290, 292, 295, 301, 303, 304, 309, 311, 341-782, 784, 785, 790, 792, 794, 796, 800, 802, 804, 807, 808, 811-826, 1240, 1243, 1247, 1269, 1272, 1280, 1283, 1285, 1286, 1289, 1300, 1309, 1318, 1319, 1327, 1335, 1339, 1346, 1359, 1369, 1370, 1371, 1393, 1398, 1405, 1408, 1413, 1414, 1417, 1422, 1429, 1432, 1435, 1436, 1438-1442, 1447, 1450, 1453, 1463, 1467, 1470, 1473, 1475, 1482, 1486, 1491-1494, 1501, 1505, 1506, 1514-1517, 1520, 1522, 1524, 1535, 1538, 1542, 1543, 1547, 1554, 1557, 1559, 1561, 1563, 1669 and 1676.

60. A diagnostic kit, comprising:

- (a) an oligonucleotide according to claim 59; and
- (b) a diagnostic reagent for use in a polymerase chain reaction or hybridization assay.

COMPOSITIONS AND METHODS FOR THE
THERAPY AND DIAGNOSIS OF LUNG CANCER

ABSTRACT OF THE DISCLOSURE

Compositions and methods for the therapy and diagnosis of cancer, such as lung cancer, are disclosed. Compositions may comprise one or more lung tumor proteins, immunogenic portions thereof, or polynucleotides that encode such portions. Alternatively, a therapeutic composition may comprise an antigen presenting cell that expresses a lung tumor protein, or a T cell that is specific for cells expressing such a protein. Such compositions may be used, for example, for the prevention and treatment of diseases such as lung cancer. Diagnostic methods based on detecting a lung tumor protein, or mRNA encoding such a protein, in a sample are also provided.

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IN THE UNITED STATES PATENT AND TRADEMARK OFFICE

Applicants : Tongtong Wang et al.
Filed : August 29, 2000
For : COMPOSITIONS AND METHODS FOR ^{THE} THERAPY AND
DIAGNOSIS OF LUNG CANCER

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Assistant Commissioner for Patents
Washington, D.C. 20231

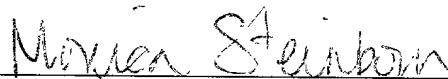
DECLARATION

Sir:

I, Monica Steinborn, in accordance with 37 C.F.R. § 1.821(f) do hereby declare that, to the best of my knowledge, the content of the paper entitled "Sequence Listing" and the computer readable copy contained within the floppy disk are the same.

I declare further that all statements made herein of my own knowledge are true and that all statements made on information and belief are believed to be true and further that these statements are made with the knowledge that willful false statements and the like so made are punishable by fine or imprisonment, or both, under Section 1001 of Title 18 of the United States Code and that such willful false statements may jeopardize the validity of the application or any patent issuing thereon.

Dated this 29th day of August, 2000.



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 ttagagttta aatttcagct tccctacttt gtaattgact gact 464

<210> 4
 <211> 510
 <212> DNA
 <213> Homo sapien

<400> 4
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 ccagttggat gtgacagaga tctttcagta taggtctaag tcaagagtag cctctgggtt 120
 gaggtgggct gggagattaa catcttacct ggggtccttc agataaacct gttggttttt 180
 cctgtctcat acaggcccat cttaagtttt gatgttgaat taaaactact tctaccccct 240
 tagttataaa aaaggccaca aggagcattt atgtggatat ctggaagtga gatagttatt 300
 ccattcccag gaaaagaaaa ataaagctaa gttacaaaac taaatctata tgcaataaag 360
 ttattatata ctgctttgtt taagcagagt cctctggaat ttatgtacag tacattagtt 420
 ttcagctatt tatattccac aagttagacc ttaagattct ctgggtttta gacaattgtt 480
 aaagatactt ctaaagctct gagcagttca 510

<210> 5
 <211> 452
 <212> DNA
 <213> Homo sapien

<400> 5
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 agtggtagat ttagaagaag agaaccataa acttttgcta gaaaatcagc ttttacgaga 180
 gaaaactcat ggccttgtag ttgagaacca ggagttaaga cagcgttggg ggatggatgc 240
 cctggttgct gaagaggagg cggaagccaa ggtaaatcat ctcttttatt tgggtgcctca 300
 tgtgagtact ggttccaagt gacatgaccc agcgattatg tttacagtct ggacttctga 360
 tcaagagcgt tcttgaaatt ttccttcagt ttaagacat tttcatgcag gcagagtgtt 420
 cttcccctaa aggcattga cactcatttt tt 452

<210> 6
 <211> 336
 <212> DNA
 <213> Homo sapien

<400> 6
 tatagagtgc tgacatctga cattgagaaa ttcatgccta ttgtttatac tcccactgtg 60

ggtctggctt	gccaacaata	tagtttggtg	tttcggaagc	caagaggtct	ctttattact	120
atccacgata	gagggcatat	tgcttcagtt	ctcaatgcat	ggccagaaga	tgatcatcaag	180
gccattgtgg	tgactgatgg	agagcgtatt	cttggttggtg	gagaccttgg	ctgtaatgga	240
atgggcatcc	ctgtgggtaa	attggctcta	tatacagctt	gcggagggat	gaatcctcaa	300
gaatgtctgc	ctgtcattct	ggatgtggga	accgaa			336

<210> 7

<211> 376

<212> DNA

<213> Homo sapien

<400> 7

ctgtgggaaa	cttcattggt	ctgtacaaag	tactagctaa	accagaaagg	tgattccagg	60
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atatctttgg	ataatgttat	ttctatTTTT	tatttttttt	cattagaagt	taccaaatta	180
agatggtaag	acctctgaga	ccaaaatttt	gtcccatctc	tacccctca	caactgctta	240
cagaatggat	catgtcccc	ttatgttgag	gtgaccactt	aattgctttc	ctgcctcctt	300
gaaagaaaga	aagaaagaag	actgtgtttt	tgccactgat	ttagccatgt	gaaactcatc	360
tcattaccct	tttctg					376

<210> 8

<211> 406

<212> DNA

<213> Homo sapien

<400> 8

ggtagggagc	aattctatta	tttggcattg	catggctggg	ttgaattaaa	acagggagtg	60
agaacaggtg	agtctagaag	tccaactctg	aaaaggacca	ctgtacattt	gaacacacgg	120
ctgtgttaaa	gatgctgcta	atgtcagtc	ctgggtgcac	taaaggatct	cttattttat	180
gtaaaacgtt	gggattgaca	agatagatct	gatactctgt	taagttacct	tctgaagcta	240
cttcttgtga	aataactaat	acagcatcat	cctgccaaagc	gaaagaggca	ggcataagca	300
aggacaaatt	aaaagggggg	aagagcctta	tcatgatgag	gagtcttggt	ttgacatctt	360
gggaaaagct	gtccatagtg	tgaagtcgtc	aatttctcac	catggt		406

<210> 9

<211> 330

<212> DNA

<213> Homo sapien

<400> 9

actactacca	agagctgcag	agagacattt	ctgaaatggt	tttgcagatt	tataaacaag	60
ggggttttct	gggcctctcc	aatattaagt	tcaggccagg	atctgtgggtg	gtacaattga	120
ctctggcctt	ccgagaagg	accatcaatg	tccacgacgt	ggagacacag	ttcaatcagt	180
ataaaacgga	agcagcctct	cgatataacc	tgacgatctc	agacgtcagc	gtgagtgatg	240
tgccatttcc	tttctctgcc	cagtctgggg	ctgggggtgcc	aggctggggc	atcgcgctgc	300
tgggtgctgg	ctgtgttctg	gttgcgctgg				330

<210> 10

<211> 449

<212> DNA

<213> Homo sapien

<400> 10

ctgacggctt	tgctgtccca	gagccgcta	aacgcaagaa	aagtcgatgg	gacagttaga	60
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ggggatgtgc taaagcgtga aatcagttgt ccttaatttt tagaaagatt ttggtaacta 120
ggtgtctcag ggctgggttg gggccaaaag tgtaaggacc ccctgccctt agtggagagc 180
tgagagcttg agacattacc ccttcacagc aaggaatttt cggatgtttt cttgggaagc 240
tgttttggtc cttggaagca gtgagagctg ggaagcttct tttggctcta ggtgagttgt 300
catgcgggta agttgaggtt atcttgggat aaagggtctt ctagggcaca aaactcactc 360
taggtttata ttgtatgtag cttatatattt ttactaaggt gtcaccttat aagcatctat 420
aaattgagtt ctttttctta gttgtatgg 449

```

```

<210> 11
<211> 472
<212> DNA
<213> Homo sapien

```

```

<400> 11
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aacccttggg ggataagaca gccacacatg gctcaggctg ttaggtgtcc actgtcacag 180
tccaaagaga aaggtagcggc ctccaagggg gcagcttaag ccaacatgta agacttgggc 240
acgatgaaaag gacggggggtc cagctacgaa tgtttttggc cttgatgtca agttgccagc 300
tactggaagg caggagcagt ttctttcttt tcccactctg tgctgggtac ttgggagagg 360
cgaaataaat accagactgt ccaactcctca gcctaaggtc cttctcaagt cctgcacact 420
cagcacttgc tctttaacgt ggcataatgt ccccatctt cccctggtta tg 472

```

```

<210> 12
<211> 371
<212> DNA
<213> Homo sapien

```

```

<400> 12
tttttttttt tttttttttt ttttggarat ttgkacatt ttattcagwa tttctgctgc 60
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gtccacaggt acctaccccc tggactgcag caactttatt accttaacta gcacaraaca 180
gaggttgatt taaactcctt acactcactt ctcaratcaa tgaatgggca aaraaacmcc 240
tcatggctct gggaaggcat gctgaracco gtttttgcaa gtccctgagga atggaaraat 300
atagctgccca ggtatcccaa gtctagggca gggagggkag tatcggcac ctttctactg 360
cattctgttg g
371

```

```

<210> 13
<211> 493
<212> DNA
<213> Homo sapien

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<220>
<221> misc_feature
<222> (1)...(493)
<223> n = A,T,C or G

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```

<400> 13
ccagtccaac ctgctcctca ttattgtata aatgagcaga atcaatatgg cggaagccag 60
ctycaattgc caatttggtg gcctctaaag ctttactttt aggaacctct gcaggcgcac 120
aggtgccaaa tcccaggaca ggcataaggt gaccatcatt cagcttcaca cactgatatt 180
tcgaatccat ttctgtcnnn nnnnnnnnnn nnnnnnnnnn nnnnnnnnnn nnnnnnnnnn 240
caacctgctc ctcatatttg taaacatgtg cagaatcaat atggcggaac ccagcttcta 300
ttgctaattt tgtgacctcc aaagctttac ttctcggaac cttgggttctt ccgagcgctc 360

```

agcaatccccg cccgagcttct ttgagacgtc ctcaggtgtc ctttgacgat gcgtccctcca 420
 ctttcacaca ctctagcatt ccttcaactgg ggtcttcatt gcccacatt gggcagccag 480
 gaatgttggg gtg 493

<210> 14
 <211> 540
 <212> DNA
 <213> Homo sapien

<400> 14
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 gtcttgact ccaggggtgga agtcatggta tagagctgag tcaactgggtc catttccttt 120
 ttaaaattat gaccaccgct ccttcaaggg gatgtagcac ttttccattc ctgtaccatg 180
 tgatattgcc atctggataa ctgtcttctg aaatgcagtc acccaacttt tttagctgct 240
 ctgtttcgag aaacagtgtc ttgcttacia tttcagggtt agatgggttg ttgaacacct 300
 tgactattgt aggtgcctca aacacgttgt cctcagttac tagcatgcac acaaactctc 360
 tttcatcact gatccttgca ttactgatag acaaagtgtg gttttctgag aggttcaatc 420
 tgtctttgta ttctggtaca tcgtcgtact gcacactttt ctttgtagag gatctgaagg 480
 caataaatac tggggagcca tcgggccttt catattttcca tttgcccata catgagattc 540

<210> 15
 <211> 421
 <212> DNA
 <213> Homo sapien

<400> 15
 taccacctc cagcctccca tgtgagcctg tccttatgta tagtgtocaa cctctgattc 60
 tagcagtcaa gtgtcttccc caatcctaata gtcccctgat atgtctctag cgaacttgacc 120
 atctcttggt ccttgggact ggggccagcc tcttgtctgc ccaactccct ctcatagtc 180
 agatagcccc aaaggctcta tctttagctc ccagagaaact ttttggctct cagtatttcc 240
 cttccctttt ccttccattt ccccaact gggggaggga agggagaaca ggggcacctg 300
 atcatcaatc tccctgccc ctctcttgaa gcccctaga tttggatgaa gagcaggcca 360
 gtgagcaggg caaagcctgc taggagcaga atgaccttga ggatcctttg ctccagaactg 420
 g 421

<210> 16
 <211> 236
 <212> DNA
 <213> Homo sapien

<400> 16
 gccgtgtgtg cttttccag tgccgaggta cctatcgctc acggccagga gcttgtcgtg 60
 gctgacagca aagagctgct ctctgtgggc ctgcttcac tcattccgaga ggccgtacaa 120
 gaagtgggtc attcctttgt ctgaaggagc gacaggagca tctacgggtg agaagacaga 180
 aagtttggct tcgtcgatgt cttgctgtgt gaattttcca gacttagccc agtoga 236

<210> 17
 <211> 424
 <212> DNA
 <213> Homo sapien

<400> 17
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 cattcagcct ttaccaatct tgtcctccaa aaaaacgaga agacatactg aaggcatgca 120

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<210> 18
<211> 154
<212> DNA
<213> Homo sapien
```

```
<210> 19
<211> 445
<212> DNA
<213> Homo sapien
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```
<210> 20
<211> 211
<212> DNA
<213> Homo sapien
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```
<210> 21
<211> 396
<212> DNA
<213> Homo sapien
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<400> 21						
tgccctgtga	ttggattgcc	acacgggtca	cattgcatgc	aagtttgctg	agctgaagga	60
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tgatgctgcc	attgttgata	tggttcctgg	caagcccatg	tgtgttgaga	gcttctcaga	180
ctatccacct	ttgggtcgct	ttgctgttcg	tgatatgaga	cagacagttg	cggtgggtgt	240
catcaaagca	gtggacaaga	aggctgctgg	agctggcaag	gtcaccaagt	ctgccagaa	300

agctcagaag gctaaatgaa tattatccct aatacctgcc accccactct taatcagtgg 360
 tggaagaacg gtctcagaac tgtttgtttc aattgg 396

<210> 22
 <211> 277
 <212> DNA
 <213> Homo sapien

<400> 22
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 aagccggcaa tctgctcccc gctgtccccg tacttccacta accagggcgg gcgctgcacc 120
 tccatcttct gggttgaggga atccacaaac cactcatccc ccatgaaatt gcaggccatg 180
 tctacatctc cattatataa taggatctgg gattttctgtg agctaagcag cttcagatac 240
 tgggagttca tgcttcggta gagacggcgg tactgta 277

<210> 23
 <211> 634
 <212> DNA
 <213> Homo sapien

<400> 23
 tctgaccatc catatccaat gttctcattt aaacattacc cagcatcatt gtttataatc 60
 agaaactctg gtccttctgt ctgggtggcacc ttagagtctt ttgtgccata atgcagcagt 120
 atggagggag gattttatgg agaaatgggg atagtcttca tgaccacaaa taaataaagg 180
 aaaactaagc tgcatgtgtg gttttgaaaa gggttattata cttcttaaca attctttttt 240
 tcagggactt ttctagctgt atgactgtta cttgaccttc tttgaaaagc attcccaaaa 300
 tgctctatct tagatagatt aacattaacc aacataatct tttttagatc gagtcagcat 360
 aaattttctaa gtcagcctct agtcgtgggt catctctttc acctgcattt tatttggtgt 420
 ttgtctgaag aaaggaaaaga ggaaagcaaaa tacgaattgt actattttgta ccaaattctt 480
 gggattcatt ggcaaataat ttcaagtgtg tgtattatta aatagaaaaa aaaaattttg 540
 tttcctaggt tgaaggtcta attgatacgt ttgacttatg atgaccattt atgcactttc 600
 aaatgaattt gctttcaaaa taaatgaaga gcag 634

<210> 24
 <211> 512
 <212> DNA
 <213> Homo sapien

<400> 24
 gcaaaacaag cctaagcaag cacaacgaag agcagaagtc agtgaaatta aaaagaggaa 60
 aaagaaaaat cataaaaaatc ataaaaagtt atttctttga aaagatcaat gaaatttagc 120
 aagactgaca cagataaaaa ggaattagac ccaaatacgt gaacaggaat gaaatagagg 180
 atatcactac agaggctgca gccattgaaa ggataattag gaaatcccac agataacttt 240
 gtgctcataa atttgacaat gtagaggaaa tatcttttagt ttttaattagc tttttatttt 300
 agtttttctc aaaaactaaa acttaataaaa actcaaccaa gacaaaatag acaatcagaa 360
 tgtaggcata cctcagagat gtggcggatt tggtttcaga ctactgcaat aaaccaata 420
 tggcaataaa aggagtcaca gaaagtgtt tcccagtgtg tatatataaa agttacattt 480
 actctatgaa gtgcaataac attttgtcta aa 512

<210> 25
 <211> 461
 <212> DNA
 <213> Homo sapien

<400> 25

ctctgtttca	gcacctcatt	gggattattg	aactcattaa	attctttaca	tgaacttgaa	60
ttgttcattg	aaatctctag	ccatttccct	ggttaaacag	gataatcttt	ttttttcact	120
aaagaacatt	cgtgggtggt	tagtgatgag	gttaatatcc	ccctcttgtc	cacctccaca	180
ttggaaaaac	cacgttggac	tgagttttga	ggagcaaaga	actaatcact	tgaccaaagg	240
ggccctgtat	ccccacaagc	cctgggtatt	tttctctcat	agagagaaga	gggtctgtat	300
ggatacctga	aaatgtgatt	ttatatattc	ttggcatcca	ggggagaaaa	atcaaaaagc	360
aaggaagtta	cagttatctc	cccagaaatt	aatgggtcat	gtcaagacta	taggttttca	420
tttcttctg	ttgcttggtt	gaatgatggt	cttgtgggaa	a		461

<210> 26

<211> 317

<212> DNA

<213> Homo sapien

<400> 26

tgctggagtc	ggaactgctg	cctttgtttg	ggggccttgt	ttcttaaadc	agttccctct	60
taggatttat	tacactaaaa	aaaaattagt	ttttgaaaag	aaataggaga	atacagaaac	120
atgaatttca	cgaggctatc	atctaacagt	gggggctttc	tacacacgtg	gtgccaaaat	180
gtgtcattct	gagtcaattg	caattcctct	ctaggagtga	aaagagataa	aagataagcc	240
aagaaccctg	gacagattct	tggtgttggt	gacaaagagg	aaaggacctg	agaatggggc	300
tggtggggag	aggggggg					317

<210> 27

<211> 250

<212> DNA

<213> Homo sapien

<400> 27

taattgctgt	gattattaga	attctatcat	gactgtattg	tagtttttgc	tctattycag	60
ataagcmaga	tctaagaagt	tatcaaaact	attctttaaa	atgctaaagc	aggtaacttt	120
ttcttccatt	attttttcct	cctaccactg	agttttgtaa	tgaattcctt	gtgtatacaa	180
gcaatacagg	tgaataactaa	actgttattt	ttagcttctt	caaaagctat	tttagaaaagc	240
ttcctggaaa						250

<210> 28

<211> 532

<212> DNA

<213> Homo sapien

<400> 28

cctatatcat	tcatttatac	agaagctgct	tgctgcttag	caagttggtg	ggtttgattt	60
tccttggttg	ctttgcagac	ctcccttgag	aggattcctt	ctggatggag	atttctttgt	120
tgctgtctcc	cttgccacaa	ctctgaccaa	gattgcattg	cgctatgtag	ctttggttca	180
ggagaagaaa	aagcaaaatt	cttttggtgc	tgaggctatg	ttgctcatgg	ctactatcct	240
gcatttgga	aaatcctctc	ttcctaagaa	gccaaattact	gatgatgatg	tggatcgaat	300
ttccctgtgc	ctcaaggtct	tgtctgaatg	ttcaccttta	atgaatgaca	ttttcaataa	360
ggaatgcaga	cagtcccttt	ctcacatggt	atctgctaaa	ctagaagaag	agaaattatc	420
ccaaaagaaa	gaatctgaaa	agaggaatgt	gacagtacag	cctgatgacc	ccatttccct	480
catgcaacta	actgctaaga	atgaaatgaa	ctgcaaggaa	gatcagtttc	ag	532

<210> 29

<211> 486

<212> DNA

<213> Homo sapien

<400> 29

ctgttttttg	acttaattaa	cywttgcaag	tggaaccaa	gaaataattg	tagcataact	60
ctctctattg	tcatgttgct	tctttctgca	aatatatctt	acaagttaga	ctttaaacct	120
ttgatctccc	acaccaaag	agaaaataat	atztatatgg	aagtaatttt	attttagtgt	180
ttgtgattta	ttgtggagag	caggbgttta	aaaatttttag	aatttccttt	taacaaaatc	240
aaatacattg	ttaaggtaac	aaagaataat	tcactatttc	agcatttcaa	agcaacatat	300
tctacaactt	caaagatatt	tgcaaaaata	atacaactgt	tgaagttcaa	atgttatgga	360
aagaaacatt	agaagtatga	aaagtggtag	aaaaacatgt	ttctttttat	tctcttggtat	420
atatatctat	atatttagga	aaatacatat	atgtatgtgt	atgtatatat	atgtatgaaa	480
atatac						486

<210> 30

<211> 240

<212> DNA

<213> Homo sapien

<400> 30

aagacctgag	gaaggaaaac	aaattggctt	cctgctgaag	aakcaaaata	gacatttttt	60
aatgtctctt	gacccagtt	ccaagtccac	cctgttgcc	gttcttctc	ccaccttttg	120
gggttctata	actgcatccc	ccacacatct	ttcaccacca	cccatacat	accagctctc	180
ctgttggtggg	attcaggaca	taggaagagt	tgctgaaggc	acgggtgctt	ttgggattcg	240

<210> 31

<211> 233

<212> DNA

<213> Homo sapien

<400> 31

ccattgatgc	aggatatcgg	cacattgact	gtgcctatgt	ctatcagaat	gaacatgaag	60
tgggggaagc	catccaagag	aagatccaag	agaaggctgt	gaagcgggag	gacctgttca	120
tcgtcagcaa	gttggtggcc	actttctttg	agagaccct	tgtgaggaaa	gcctttgaga	180
agacctcaa	ggacctgaag	ctgagctatc	tggacgtcta	tcttattcac	tgg	233

<210> 32

<211> 233

<212> DNA

<213> Homo sapien

<400> 32

gaggaatgct	ggactggagg	cccctggagc	cagatggcaa	gagggtgaca	gcttcctttc	60
ctgtgtgtac	tctgtccagt	tccttttagaa	aaaatggatg	cccagaggac	tcccaaccct	120
ggcttgggg	caagaaacag	ccagcaagag	ttaggggctt	tagggcactg	ggctgttgtt	180
ccattgaagc	cgactctggc	cctggccctt	acttgcttct	ctagctctct	agg	233

<210> 33

<211> 319

<212> DNA

<213> Homo sapien

<400> 33

ctgggcctgg	atggtotagg	atagccttac	tcacttgctt	ggcaggtgac	aggctgttgg	60
ctggaattgc	ttggttctcc	tccatgtggc	ctctccagta	ggctagctca	ggcttattca	120

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catgatggct tcaggattcc aaagagagtg agagtagaag ctgaaagact tcttgagttc 180
ttggcctgga actgggacta ggacagtgtc acttctgcta agttcttttg gtcagagcaa 240
atcacaaggc tttacccaga ttcaagggat gaaaaacaga ctacatgtct tgatgagggg 300
aaccacaaag agcttgtgg 319

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```

<210> 34
<211> 340
<212> DNA
<213> Homo sapien

```

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<400> 34
tacagattta attcatgtta ttaactccct gccttttacc tcctccctcc tcccttggca 60
caactgccag atggatgtgg ctggaagtcaggagacattc tcgtgggttc gtgggcctag 120
ggtacaaatg acctcagcgt gacagcaaac aggacagaga agaccaggct cttactcagg 180
aatccaccag ccaggagaat gacaatgttg aacaccggaa ccctgatgat atctgtcaca 240
tttgtaaggt tgatttcaga gtcaggagtg gagacatcgg cagttgactt ggggtggagct 300
tgggtcacag ttctggggct ggtatagagt gggcacaagg 340

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<210> 35
<211> 170
<212> DNA
<213> Homo sapien

```

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<400> 35
acatgggtcc ttcactcctc gctgagatgt tgccggcagcc ttttcttcca atgcgggttg 60
ggcaggagaa tccacgatg taatgttttc acctttttcc ctgaggggtgc tttctgagga 120
accagycctt aagaggtggg gtcttggatt cctgaccacag gcgtccggca 170

```

```

<210> 36
<211> 475
<212> DNA
<213> Homo sapien

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```

<400> 36
ctgttttttg acttaattaa ccattgcaag tggaaaccaa gaaataattg tagcataact 60
ctctctattg kcatgttgct tctttctgca aatatatctt agaagttaga ctttaaacct 120
ttgatctccc acacaaaag agaaaataat atttatatgg aagtaatttt attttagtgt 180
ttgtgattta ttgtggagag caggtgttta aaaatttttag aatttcttta acaaaattct 240
aaagagaaaa taaaaaagaa atcacagtat ttacagagat aacagaatgg cttagccatg 300
caaaacaaat aactttgggt tttccccttt tactttgggt taaatgttga ccaagattca 360
atTTTTTTTt ctgccaaata aaacttcaat aaaagttagt aggcaaaata acgtattttc 420
TTTTTTTccc ataattttt atacagcatc gagtctaaga atatttttatg cttt 475

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<210> 37
<211> 246
<212> DNA
<213> Homo sapien

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<400> 37
ccttgagctt gggccgggca ctgaggcgcc ccacatatgc tgagagcagg gggaaacgcat 60
ccaggcagcc aggggctagg acctcatgga tcagcagcaa gtccagcagg ttgtagtcag 120
cgaaggagat ctggtctccc acaatgaagg tcttgctccc ctggttctgg gacagcaggg 180
tctcaaaagg cttcagttgc cggggcagtg ctttcacata gtcaccttg cccacctcat 240
agttgg 246

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<210> 38
 <211> 512
 <212> DNA
 <213> Homo sapien

<400> 38
 gctggaagtg aaatgcagat cagacccatt gtgatgtcac agaaagatgg ggacaggcca 60
 aagaaaaaag tgactttcaa ctcttcttcc atcattttta tcatcaccag tgatgaatca 120
 ctgtcagttg acgacagcga caaaaccaat ggggtccaaag ttgatgtaat ccaagttcgt 180
 cctttgtagg aatgaagaat ggcaacgaaa gatggggcct taaattggat gccacttttg 240
 gactttcatc ataagaagtg tctggaatac cgtttctatg taatatcaac agaaccttgt 300
 ggtccagcag gaaatccgaa ttgccatat gctcttgggc ctccaggaaga ggttgaacaa 360
 aaacaaattc ttttaattca acgggtgctt tacataatga aaaaaccact tgtggcacac 420
 gatgggcac taacatcatc atcttctaata gtgttggaga ttttcatttc aaatatattt 480
 tttaaattac tctattttcc aaaacacgta at 512

<210> 39
 <211> 370
 <212> DNA
 <213> Homo sapien

<400> 39
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 atgtactcga ctctgtccta tttagccttc ccataacctga cttctaataca cttttcctgg 120
 tgccctycca tctccctaac cccccctcac agggatgcct cctcccaagg ctccagaaac 180
 tctgacctc gcaactgctgg agggagccca tgaattgctg gtcaatatcg ctcatcctct 240
 akactccatc ctgctgtgct ttcttcctac aagagctaga gaggcactga ctgataaata 300
 cctgtcacct gcccctttcc cagaggggtga aactccaccc actcccaactg cagaaatgaa 360
 tcttaaattg 370

<210> 40
 <211> 204
 <212> DNA
 <213> Homo sapien

<400> 40
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 ggagcagagc agacctgtt tttagtgggt ccatgggata aaatgggatt ggaggagcta 120
 gaagaattca gggctctggc caatctgcca gtcttcctga aatatcgaaa atacaccagg 180
 gctgctatat cagagccacc ctgg 204

<210> 41
 <211> 447
 <212> DNA
 <213> Homo sapien

<400> 41
 caggcagcaa ttcgtaaaga attaaatgag taaaaaagta atgaaatgga ggtacatgca 60
 tcaagcaagc acttgacaag attccacagg ccatagagat tttcttctga gaagaatttg 120
 tgtttaattt tttgatacca aactgaaca ttcacaggg aactttctctg aagttcagct 180
 caagactacc ctacctgctg tgtttgtgag aagagtagga tcacacacac aggtgcaatc 240
 ttgaccacac ttacctgcaa gaggagtaac cagaggacac acttccttcc ttctttggtg 300
 tctgaggagt gtgaactgtt ggggtcagtt aagacccaac ataactctat cagaagaaaa 360

ctgttgtttg cttttcaacc ttgttttaca gttctgcagt gtagtggagg acgggcaacg 420
 tgcattgtgca ggctcaccac tcccagg 447

<210> 42
 <211> 498
 <212> DNA
 <213> Homo sapien

<400> 42
 ctggttttgt aaaaacagtc tctttattct actgtgctga aacctcacc aatatagaaa 60
 attagattct cattgcactg aactatatatt atatgcctaa gtatgtagaa gtaaaattat 120
 ataccccaaa aggattttat cttgttgtat atattaaatg ttatttctgc atatagggtc 180
 ttttatggag aaactgatga tgataagctt aatactcact tgtttagcag catctgaatg 240
 cacaaatgct ttatatatct cttctgcttt acagggcaaa agatcagact ctgttttctt 300
 atagtcttca caagccagcc agaactcaat attctcctca ctgaattcag actttaggaa 360
 acttccaaag acattttgac cagtttggtt ggcaagaagt tttccagag attgagacca 420
 ttgcattact tcagcagcag aaagtacatc cttggacttg gaagatttca ttccagattc 480
 cagatgtggg atcataga 498

<210> 43
 <211> 312
 <212> DNA
 <213> Homo sapien

<400> 43
 caggaaggcg gccagaatg tgagtgcaaa gattggttcc tgagagcccc gagaagaaaa 60
 ttcattgacag tgtctgggct gccaaagaag cagtgcacct gtgatcattt caagggcaat 120
 gtgaagaaaa caagacacca aaggcaccac agaaagccaa acaagcattc cagagcctgc 180
 cagcaatttc tcaaacaatg tcagctaaga agctttgtct tgcccttgta ggagctctga 240
 gcgcccactc ttccaattaa acattctcag ccaagaagac agtgagcaca cctaccagac 300
 actctttctt tc 312

<210> 44
 <211> 417
 <212> DNA
 <213> Homo sapien

<400> 44
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 ccttctcaag ccatgtctca gagctgagag gcatcccagc aagttttgca gctcacagtt 120
 ttttccgtaa attacttatt ctataaaatt ggagtaggcc ataaactttg gagggcccta 180
 gaccaatttt ttggattatt tttcgtcttc tatcattccg ctgatcttag atattctctg 240
 cattaaatat taaatatcac ttctaggttg aaaaatcccc ctaaaaatat ttctagctca 300
 gatttttctt ccaaattctg caatagaaga tcacaatgtg aactctgcat ctccatgtta 360
 aagtctaattg gacattcaca cttagcatgt ctcaaagaaa tctcatgtaa accatgg 417

<210> 45
 <211> 494
 <212> DNA
 <213> Homo sapien

<400> 45
 cgcggtgtctg tgggtatgtg acacgtgcat gttctgcatg tctgtaggtc acacatgctt 60
 tgggtgcatgt acacgtgtgt gtgtgtatgc gtgtaggagc tcacacttgt gtacacgttt 120

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gtgtgcatgc atgtgtgcag gagcttgca c gtttgtggtg ggtacatgta catatgtgag 180
tgatcctgtg tgcaagcccc catgtggaca tggctatgag tgagcgtgga gccaaaagcc 240
aggtaacacg catgcagcag gccactgtg cgtgtctgag acggtctgtg gcagggactg 300
ggtgtgaatc atgcagcagg cccactgtgc gtgtctgaga cggctctgtg cagggactgg 360
gtgtgaatca gtgaccgtgt ctctgaccaa catgctgaat tacaaattga taatttatta 420
acctgtgcag caacaaataa gatttttcaa aactcaacaa agtgctcaaa gttgacatta 480
cttgcttcaa agtt 494

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<210> 46
<211> 516
<212> DNA
<213> Homo sapien

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<400> 46
ccagtccaac ctgctcctca ttattgtata aatgagcaga atctatatgg cggaacccag 60
cttctattgc taattttgtg acctocaaag ctttacttct cggaacctcc tcctttggcc 120
gtcatttgat cattcaactc tttgtcagtg gcaactcccg ctattttggt gtgttggttt 180
gttactacac agtgagcaca aacatggtgg tccaatacag aggcctcttc tgtcagggtg 240
caaccagaaa gttcatctaa cactgtgata tttgcatact tcttgaacag ttgttggtctg 300
aagattcatt tgatgaatcg atttttcaaa agagatgatt cttggttctt ccgagcgctc 360
agctctcccg ccgagcttct ttgagacgtc ctcagggtgtc ctttgacgat gcgtcctcca 420
ctttcacaca ctctagcatt ccttacttgg ggtcttcatt gcccacatt gggcagccag 480
gaatgttggg gtgatcagac acaacaccag gtcatg 516

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<210> 47
<211> 459
<212> DNA
<213> Homo sapien

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<400> 47
ccaattcaga gtggcattct gcatttctgt ggcttccaag tcttagaacc tcaactgaca 60
tatagcattg ggcacactcc agcagacgcc cgaattcaaa tcttggaagg atggaagaaa 120
cgcctggaga atatttgga tgagacacca ctgtattttg ctccaagcag cctctttgac 180
ctaaacttcc aggcaggatt cttaatgaaa aaagaggtag aggatgagga gaaaaacaag 240
aaatttgccc tttctgtggg ccatacattg ggcaagtcca tcccaactga caaccagatc 300
aaagctagaa aatgagattc cttagcctgg atttcttct aacatgttat caaatctggg 360
tatctttcca ggcttccctg acttgcttta gtttttaaga tttgtgtttt tctttttcca 420
caaggaataa atgagaggga atcgaksaaa aaaaaaaaaa 459

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<210> 48
<211> 430
<212> DNA
<213> Homo sapien

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<400> 48
cctatattca gccacagcct ctgggagtgg tgctgataat cggagcttgg aattaccctt 60
tcgtttcac cattcagcca ctgataggag ccatcgctgc aggaaatgct gtgattataa 120
agccttctga actgagtga aatacagcca agatcttggc aaagcttctc cctcagtatt 180
tagaccagga tctctatatt gttattaatg gtgggtgtga ggaaaccacg gagctcctga 240
agcagcgatt tgaccacatt ttctatacgg gaaacactgc ggttggcaaa attgtcatgg 300
aagctgctgc caagcatctg acccctgtga ctcttgaact gggagggaaa agtccatgtt 360
atattgataa agattgtgac ctggacattg tttgcagacg cataacctgg ggaaaataca 420
tgaattgtgg 430

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<210> 49
 <211> 288
 <212> DNA
 <213> Homo sapien

<400> 49
 ccatccgaag caagattkca gatggcagtg tgaagagaga agacatattc tacacttcaa 60
 agctttggwg caattcccat cgaccagagt tgggccgacc agccttggaa aggtcactga 120
 aaaatcttca attggattat gttgacctct accttattca ttttccagtg tctgtaaagc 180
 caggtgagga agtgatccca aaagatgaaa atggaaaaat actatttgac acagtggatc 240
 tctgtgccac gtgggaggcc rtggagaagt gtaaagatgc aggattgg 288

<210> 50
 <211> 411
 <212> DNA
 <213> Homo sapien

<400> 50
 ccagagaatg acattcatgt ccccgaggat cccttgacaga gagtacatgg agccactgcc 60
 accagtggtg atggaaagca ctgtcttctt actccggaag ggctccttgg catacatggc 120
 agcgtaagt taagcaaact ctctatgaa cactcgtcga aaccagcctt tcagaatggc 180
 agggactcca aaccactgca gggggaactg gaatatcaca aggtctgcgg ctccagctt 240
 cttttgttca gccacaatat ctgggctcag atggccttct ttataagcca gaacagactc 300
 ggcaggatac tgaaagtctg cagggctcct cagtttacct gtgatgtcct ttctggaaat 360
 gatgggattg aagttcatgg catagaggct cgactccacc acctccatc c 411

<210> 51
 <211> 503
 <212> DNA
 <213> Homo sapien

<400> 51
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 ttgtgcaccc tccacaaaac atacaaagtt taaaagtttg gatctttttc tcagcaggta 120
 tcagttgtaa ataataaatt aggggccaaa atgcaaaacg aaaaatgaag cagctacatg 180
 tagttagtaa tttctagttt gaactgtaat tgaatattgt ggcttcatat gtattatattt 240
 atattgtact tttttcatta ttgatggttt ggactttaat aagagaaatt ccatagtttt 300
 taatatccca gaagtgaagc aatttgaaca gtgtattcta gaaaacaata cactaactga 360
 acagaagtga atgcttatat atattatgat agccttaaac ctttttcctc taatgcctta 420
 actgtcaaat aattataacc ttttaaagca taggactata gtcagcatgc tagactgaga 480
 ggtaaacact gatgcaatta aga 503

<210> 52
 <211> 503
 <212> DNA
 <213> Homo sapien

<400> 52
 gatatcttat gattaataaac aaattaaatt ttaaaacacc tgaagatata ttagaagaaa 60
 ttgtgcaccc tccacaaaac atacaaagtt taaaagtttg gatctttttc tcagcaggta 120
 tcagttgtaa ataataaatt aggggccaaa atgcaaaacg aaaaatgaag cagctacatg 180
 tagttagtaa tttctagttt gaactgtaat tgaatattgt ggcttcatat gtattatattt 240
 atattgtact tttttcatta ttgatggttt ggactttaat aagagaaatt ccatagtttt 300
 taatatccca gaagtgaagc aatttgaaca gtgtattcta gaaaacaata cactaactga 360

acagaagtga atgcttatat atattatgat agccttaaac ctttttcctc taatgcctta 420
actgtcaaat aattataacc ttttaaagca taggactata gtcagcatgc tagactgaga 480
ggtaaact gatgcaatta aga 503

<210> 53
<211> 531
<212> DNA
<213> Homo sapien

<400> 53
tttttttttt tttttaaaat gaggatattt tattatttca ggtaattttc ccagaggkga 60
gaatagtaca tgggaaattc tctttaggcc aggtctagta ttacagkgtg gkgctcaagg 120
ccgcccatac gaacagtgat actctcccaa cagatttcat ccaccccgtc tccactaact 180
tttgccataa aaattcctct gaattgtatc ttcttggaag aagtaaataat ctgttcgact 240
atacaaagaa acagagaaac cactcccatt gcaatcaatc ttcaagagag ggagcaggca 300
agccgtgttc tttctgctga gttttataga ctctgacaag ctgtgaaata aacataaaca 360
gaagacaaaa cagtgcacaa aataagcagt agatgaccct gtgacaagac ggcattgcag 420
aacaagact gacgtttaaa ggggagtcac gcagagtaac atgggaacac aagcctgaca 480
acctggtcag cttccactta ctctagctcc tttgaactct caacactaaa a 531

<210> 54
<211> 450
<212> DNA
<213> Homo sapien

<400> 54
ccatgggtgt ctggagcwcc ctgaaactgt atcaaagttg tacatatttc caaacatttt 60
taaaatgaaa aggcactctc gtgttctcct cactctgtgc actttgctgt tgggtgtgaca 120
aggcatttaa agatgtttct ggcattttct ttttatttgt aagggtggtg taactatggt 180
tattggctag aaatcctgag ttttcaactg tatatatcta tagtttgtaa aaagaacaaa 240
acaaccgaga caaaccttg atgctccttg ctggcgcttg aggcctgtgg gaagatgcct 300
tttgggagag gctgtagctc agggcggtgca ctgtgaggct ggacctgttg actctgcagg 360
gggcatccat ttagcttcag gttgtcttgt ttctgtatat agtgacatag cattctgctg 420
ccatcttagc tgtggacaaa ggggggtcag 450

<210> 55
<211> 648
<212> DNA
<213> Homo sapien

<400> 55
caacttcaac cacaggctgc tggasatgat cctcarcaag ccagggtcctca agtacaagcc 60
tgtctgcaac caggtggaat gtcataccta cttcaaccag agaaaaactgc tggatttctg 120
caagtcaaaa gacattgttc tggttgecta tagtgctctg ggatcccacc gagaagaacc 180
atgggtggac ccgaactccc cgggtgctctt ggaggaccca gtcctttgtg ccttggaaca 240
aaagcacaag cgaacccag ccctgattgc cctgcgctac cagctrcagc gtgggggtgt 300
ggtcctggcc aagagctaca atgagcagcg catcagacag aacgtgcagg tgtttgaatt 360
ccagttgact tcagaggaga tgaaagccat agatggccta aacagaaatg tgcgatattt 420
gacccttgat atttttgctg gccccctaa ttatccattt tctgatgaat attaacatgg 480
agggcattgc atgaggtctg ccagaaggcc ctgcgtgtgg atggtgacac agaggatggc 540
tctatgctgg tgaactggaca catcgctctt ggtaaactct ctctgcttg gygayttcag 600
caagctacag caaagcccat tggccggaat aaatatcaag ggtcaaat 648

<210> 56

<211> 536
 <212> DNA
 <213> Homo sapien

<400> 56
 ctggcatgag aatatttttt tttttaagtg cggtagtttt taaactgttt gtttttaaac 60
 aaactataga actcttcatt gtcagcaaag caaagagtca ctgcatcaat gaaagttcaa 120
 gaacctcctg tacttaaaca cgattcgcaa cgttctgtta ttttttttgt atgttttagaa 180
 tgctgaaatg tttttgaagt taaataaaca gtattacatt tttaaaactc ttctctatta 240
 taacagtcaa tttctgactc acagcagtga acaaaccccc actccattgt atttggagac 300
 tggcctccct ataaatgtgg tagcttcttt tattactcag tggacctgcc cgggcggccg 360
 ctggaagccg aattccagca cactggcggc cgttactagt ggatccgagc tcggtaccaa 420
 gcttggccgt aatcatggtc atagctgttt cctgtgtgaa attgttatcc gtcacaatt 480
 ccacacaaca tacgagccgg aagcataaag tgtaaagcct ggggtgccta atgagt 536

<210> 57
 <211> 391
 <212> DNA
 <213> Homo sapien

<400> 57
 aggaactact gtcccagagc tgaggcaagg ggattttctca ggtcatttgg agaacaagtg 60
 ctttagtagt agtttaaagt agtaactgct actgtattta gtgggggtgga attcagaaga 120
 aatttgaaga ccagatcatg ggtggtctgc atgtgaatga acaggaatga gccggacagc 180
 ctggctgtca ttgctttctt cctccccatt tggacccttc tctgccctta catttttgtt 240
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 gaaaatttat tgtttattgg catgtgacct ttgactgatg gcttcattag cattytgttt 360
 ttcttttttg atccttaata gaaaactcaa t 391

<210> 58
 <211> 455
 <212> DNA
 <213> Homo sapien

<400> 58
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 gccctagcca acgccgcatg agagggagtg tgccgagggc ttctgagaag gtttctctca 120
 catctagaaa gaagcgctta agatgtggca gccctcttc ttcaagtggc tcttgtcctg 180
 ttgccctggg agttctcaaa ttgctgcagc agcctccacc cagcctgagg atgacatcaa 240
 tacacagagg aagaagagtc aggaaaagat gagagaagtt acagactctc ctgggcgacc 300
 ccgagagctt accattcctc agacttcttc acatggtgct aacagatttg ttcctaaaag 360
 taaagctcta gaggccgtca aattggcaat agaagccggg ttccaccata ttgattctgc 420
 acatgtttac aataatgagg agcaggttgg actgg 455

<210> 59
 <211> 398
 <212> DNA
 <213> Homo sapien

<220>
 <221> misc_feature
 <222> (1)...(398)
 <223> n = A,T,C or G

<400> 59

ctcagaggca	gcgtgcgggt	gtgctctttg	tgaaattcca	ccatggcgta	ccgtggccag	60
ggtcagaaag	tgcagaaggt	tatgggtgcag	cccatcaacc	tcctcttcag	atacttacia	120
aatagatcgc	ggattcaggt	gtggctctat	gagcaagtga	atatgcggat	agaaggctgt	180
atcattgggt	ttgatgagta	tatgaacctt	gtattagatg	atgcagaaga	gattcattct	240
aaaacaaagt	caagaaaaca	actngntcgg	atcatgctaa	aaggagataa	tattactctg	300
ctacaaagtg	tctccaacta	gaaatgatca	atgaagtgag	aaattgttga	gaaggataca	360
gtttgttttt	agatgtcctt	tgtccaatgt	gaacattt			398

<210> 60

<211> 532

<212> DNA

<213> Homo sapien

<400> 60

gacttctgag	acctggggca	cccgggcctt	tgcggcagct	actggcaggg	cctggccacc	60
tcataggact	cagttccctt	ctgaacactc	gggggacatg	ggcctctaac	tgcccactct	120
gatatgcctg	ggtgagccta	ggagggaagg	ctctgatttg	gatttctcca	gtcaaagctc	180
acagaaaaaa	acctggcact	ttgattttca	tgggatggtc	ctaacagggg	cagtcacctc	240
cgagcagttt	gggaacccag	tttcttgtcc	tgggcccctc	ggtcagcctg	gctgaattag	300
gaccttccct	tggcacaggg	gtgagaaaga	gcttggggaa	cgcttggcat	tatggagggc	360
tgggaagggc	tcaaccccga	tttggagaga	agtttgggat	ggagtgggcg	agagattgag	420
agagcgagca	ggaaaagagg	tcttggagcc	tgggactgat	ggtggataag	gcctggaaag	480
aasatgacsa	ggaggaggag	agaggggaagt	gggtggatga	ggagcaggct	ga	532

<210> 61

<211> 466

<212> DNA

<213> Homo sapien

<400> 61

gcgacggcga	cgtctctttt	gactaaaaga	cagtgtccag	tgctccagcc	taggagtcta	60
cggggaccgc	ctcccgcgcc	gccacatgc	ccaacttctc	tggcaactgg	aaaatcatcc	120
gatcggaaaa	cttcgaggaa	ttgctcaaa	tgctgggggt	gaatgtgatg	ctgaggaaga	180
ttgctgtggc	tgcagcgtcc	aagccagcag	tggagatcaa	acaggaggga	gacactttct	240
acatcaaaac	ctccaccacc	gtgcgcacca	cagagattaa	cttcaagggt	ggggaggagt	300
ttgaggagca	gactgtggat	gggaggccct	gtaagagcct	ggtgaaatgg	gagagtgaga	360
ataaaatggg	ctgtgagcag	aagctcctga	agggagaggg	ccccaagacc	togtggacca	420
gagaactgac	caacgatggg	gaactgatcc	tgaccatgac	ggcgga		466

<210> 62

<211> 548

<212> DNA

<213> Homo sapien

<400> 62

ttttgaattt	acaccaagaa	cttctcaata	aaagaaaatc	atgaatgctc	cacaatttca	60
acataccaca	agagaagtta	atttcttaac	attgtgttct	atgattattt	gtaagacctt	120
caccaagttc	tgatatcttt	taaagacata	gttcaaaatt	gcttttgaaa	atctgtattc	180
ttgaaaatat	ccttgttgtg	tattaggttt	ttaaatacca	gctaaaggat	tacctcactg	240
agtcacagct	accctcctat	tcagctcccc	aagatgatgt	gtttttgctt	accctaagag	300
aggttttctt	cttattttta	gataattcaa	gtgcttagat	aaattatgtt	ttctttaagt	360
gtttatggta	aactctttta	aagaaaattt	aatatgttat	agctgaatct	ttttggtaac	420
tttaaattct	tatcatagac	tctgtacata	tgttcaaatt	agctgcttgc	ctgatgtgtg	480

tatcatcggg gggatgacag aacaaacata tttatgatca tgaataatgt gctttgtaaa 540
aagatttc 548

<210> 63
<211> 547
<212> DNA
<213> Homo sapien

<400> 63
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cctatgtaag gccatgtgcc ccttgcccta acaactcact gcagtgtctt tcatagacac 120
atcttgacgc atttttctta aggtatgct tcagtttttc tttgtaagcc atcacaagcc 180
atagtggtag gtttgccctt tggtagagaa ggtgagttaa agctgggtgga aaaggcttat 240
tgcattgcat tcagagtaac ctgtgtgcat actctagaag agtagggaaa ataatgcttg 300
ttacaattcg acctaatatg tgcattgtaa aataaatgcc atatttcaaa caaaacacgt 360
aattttttta cagtatgttt tattaccttt tgatatctgt tgttgcaatg ttagtgatgt 420
tttaaaatgt gatcgaaaat ataatgcttc taagaaggaa cagtagtgga atgaatgtct 480
aaaagatctt tatgtgttta tggctctgcag aaggattttt gtgatgaaag gggatttttt 540
gaaaaat 547

<210> 64
<211> 528
<212> DNA
<213> Homo sapien

<220>
<221> misc_feature
<222> (1)...(528)
<223> n = A,T,C or G

<400> 64
cacctmctcc cscwggcgcc ttwctcsgac gccttgccca scggggccgcc cgacccccctg 60
srccatggac cccgctcgcc csctggggmt gtygatktct ctgcttttcc tgrckgaggc 120
tgcaactggc gatgctgatc argagccaac aggaaataac rcggagatct gkctcctgcc 180
cctagactac kgacccctgcc kggccctact tytccgytac tactacgaca ggyacacgca 240
gagctgccgc cwgttctctgk rckggggctg crasggcaac rccaacwatt yctacacckg 300
kgaggmtrc gackatgctw gstggargat agaaaaagtt cccaaasttt gccggctgma 360
agtgaatgag gacnaccagg gtgaggggta cacagataag tatttcttta atctaakkwc 420
catgacatgw gaaaaattct ttnncgggtg gngtcaccgg accggattga gaacangttt 480
gcagatgang ctactgggat gggctcctgc rcacnaaaga aantatca 528

<210> 65
<211> 547
<212> DNA
<213> Homo sapien

<220>
<221> misc_feature
<222> (1)...(547)
<223> n = A,T,C or G

<400> 65
kgaatgaasa acgaacgctg gaagtagaaa tagagcctgg ggtgagagac ggcattggagt 60
acccctttat tggagaaggt gagcctcacg tggatgggga gcctggagat ttacggttcc 120

gaatcaaagt	tgtcaagcac	ccaatatttg	aaaggagagg	agatgatttg	tacacaaatg	180
tgacagtctc	attagttgag	tcaactggtg	gctttgagat	ggatattact	cacttggatg	240
gtcacaaggt	acatatttcc	cgggataaga	tcaccaggcc	aggagcgaag	ctatggaaga	300
aaggggaagg	gctccccaac	tttgacaaca	acaatatcaa	gggctctttg	ataatcactt	360
ttgatgtgga	ttttccaaaa	gaacagttaa	cagaggaagc	gagagaangt	atcaaacagc	420
tactgaaaca	agggtcagtg	cagaagggtat	acaatggact	gcaaggatat	tgagagtgaa	480
taaaattgga	ctttgtttta	aataaagtga	ataagcgata	tttattatct	gcaagggttt	540
ttttgtg						547

<210> 66
 <211> 535
 <212> DNA
 <213> Homo sapien

<400> 66						
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gcaccgcgcc	ctccaccgcc	ggttggtggc	ctgcgtgaca	gtttcctccc	gtcgacatcg	120
aaaggaagcc	ggacgtgggc	gggcagagag	cttcacgcga	gtaggaatgg	cagcccatc	180
tatgaaggaa	agacaggtct	gctggggggc	ccgggatgag	tactggaagt	gtttagatga	240
gaacttagag	gatgcttctc	aatgcaagaa	gttaagaagc	tctttcgaat	caagttgtcc	300
ccaacagtgg	ataaaaatatt	ttgataaaaag	aagagactac	ttaaaattca	aagaaaaatt	360
tgaagcagga	caatttgagc	cttcagaaac	aactgcaaaa	tcctaggctg	ttcataaaga	420
ttgaaagtat	tctttctgga	cattgaaaaa	gctccactga	ctatggaaca	gtaatagtgt	480
gaatcatagt	gaacatcaat	acttgttccc	tatatacgac	acttgataat	taaga	535

<210> 67
 <211> 527
 <212> DNA
 <213> Homo sapien

<400> 67						
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ttcatcttct	acaaggccct	cttagctcta	aaacttgaca	gtggaataag	gaaatgtttt	120
tccaaatctg	cattgcccgt	gagatcctca	acatcagcat	gttgagatgg	acctcaaccc	180
cacctctaac	cctgaaacac	actactcgat	attatcttag	gtatgtttta	gggttttagtt	240
tgtaaaataa	taatttat	ttgaaggaaa	tataaaatat	taaagagtaa	taatagctat	300
cattttttta	gattcaatct	aaaacaatgg	actctttttt	tttccatttg	tgatgtagat	360
aagcaagaca	attttgatca	tgagtgggtga	aaagaggatc	aaacttgact	attcttgcaa	420
tggcagtcca	gcaacaagcc	tttcatttac	attaaattat	aacttttcat	tcattcctaa	480
accaaactta	aaattctgct	ttcctttgag	tagaagggtat	ttaactt		527

<210> 68
 <211> 431
 <212> DNA
 <213> Homo sapien

<400> 68						
gggaaacttc	atgggtttcc	tcatctgtca	tgctgatgat	tatatatgga	tacatttaca	60
aaaataaaaa	gcgggaattt	tcccttcgct	tgaatattat	ccctgtatat	tgcatgaatg	120
agagatttcc	catattttcca	tcagagtaat	aaatataact	gctttaattc	ttaagcataa	180
gtaaacaatga	tataaaaaata	tatgctgaat	tacttggtga	gaatgcattt	aaagctat	240
taaatgtgtt	tttattttgta	agacattact	tattaagaaa	ttggttatta	tgcttactgt	300
tctaactctgg	tggtaaagggt	attcttaaga	atttgcagggt	actacagatt	ttcaaaactg	360
aatgagagaa	aattgtataa	ccatcctgct	gwtcccttag	tgcaatacaa	taaaactctg	420

aaattaaaaac t

431

<210> 69

<211> 399

<212> DNA

<213> Homo sapien

<400> 69

gacacggcgg	acacacacaa	acacagaacc	acacagccag	tcccaggagc	ccagtaatgg	60
agagcccaa	aaagaagaac	cagcagctga	aagtccggat	cctacacctg	ggcagcagac	120
agaagaagat	caggatacag	ctgagatccc	agtgcgcgac	atggaagggtg	atctgcaaga	180
gctgcatcag	tcaaaccaccg	gggataaatc	tggattttggg	ttccggcgctc	aagggtgaaga	240
taatacctaa	agaggaacac	tgtaaaatgc	cagaagcagg	tgaagagcaa	ccacaagttt	300
aatgaagac	aagctgaaac	aacgcaagct	ggttttatat	tagatatattg	acttaaacta	360
tctcaataaa	gttttgcagc	tttcaccaar	aaaaaaaaa			399

<210> 70

<211> 479

<212> DNA

<213> Homo sapien

<400> 70

cgcggcggag	ctgtgagccg	gcgactcggg	tccctgaggt	ctggattctt	tctccgctac	60
tgagacacgg	cggacacaca	caaacacaga	accacacagc	cagtcccagg	agcccagtaa	120
tggagagccc	caaaaagaag	aaccagcagc	tgaaagtccg	gatacctaac	ctgggcagca	180
gacagaagaa	gatcaggata	cagctgagat	cccagggtgct	gggaagggaa	atgcgcgaca	240
tggaagggtga	tctgcaagag	ctgcatcagt	caaacaccgg	ggataaatct	ggattttgggt	300
tccggcgctca	aggtgaagat	aatacctaaa	gaggaacact	gtaaaatgcc	agaagcagggt	360
gaagagcaac	cacaagttta	aatgaagaca	agctgaaaca	acgcaagctg	gttttatatt	420
aggatatattg	acttaaacta	tctcaataaa	gttttgcagc	tttcaccaa	aaaaaaaaa	479

<210> 71

<211> 437

<212> DNA

<213> Homo sapien

<400> 71

ctcagcgggt	gccaacagat	catgagccat	cagctcctct	ggggccagct	ataggacaac	60
agaactctca	ccaaaggacc	agacacagtg	rgcaccatgg	gacagtgtcg	gtcagccaac	120
gcagaggatg	ctcaggaatt	cagtgatgtg	gagagggcca	ttgagaccct	catcaagaac	180
tttcaccagt	actccgtgga	gggtgggaag	gagacgctga	ccccttctga	gctacggggac	240
ctggtcaccc	agcagctgcc	ccatctcatg	ccgagcaact	gtggcctgga	agagaaaatt	300
gccaacctgg	gcagctgcaa	tgactctaaa	ctggagttca	ggagtttctg	ggagctgatt	360
ggagaagcgg	ccaagagtgt	gaagctggag	aggcctgtcc	gggggcaactg	agaactccct	420
ctggaattct	tggggggg					437

<210> 72

<211> 561

<212> DNA

<213> Homo sapien

<400> 72

ggatgggtata	ctgtaaattc	agcatatgga	gataaccatta	tcataccttg	ccgacttgac	60
gtacctcaga	atctcatgtt	tggcaaatgg	aaatatgaaa	agcccgatgg	ctccccagta	120

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<210> 73
<211> 916
<212> DNA
<213> Homo sapien
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<210> 74
<211> 547
<212> DNA
<213> Homo sapien
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<210> 75
<211> 793
<212> DNA
<213> Homo sapien
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<400> 75

tgaggaagtt	gcaagccaac	aaaaaagttc	aaggatctag	aagacgatta	aggggaaggtc	60
gttctcagt	aaaatccaaa	aaccagaaaa	aatgttttat	acaaccctaa	gtcaataacc	120
tgaccttaga	aaattgtgag	agccaagtgt	acttcaggaa	ctgaaacatc	agcaciaaaga	180
agcaatcatc	aaataattct	gaacacaaat	ttaatatatt	tttttctgaa	tgagaaacat	240
gagggaaatt	gtggagttag	cctcctgtgg	agttagcctc	ctgtggtaaa	ggaattgaag	300
aaaatataac	accttacacc	ctttttcatc	ttgacattaa	aagttctggc	taactttgga	360
atccattaga	gaaaaatcct	tgtcaccaga	ttcattacaa	ttcaaatacg	agagttgtga	420
actgttatcc	cattgaaaag	accgagcctt	gtatgtatgt	tatggatata	taaaatgcac	480
gcaagccatt	atctctccat	gggaagctaa	gttataaaaa	taggtgcttg	gtgtacaaaa	540
ctttttatat	caaaaaggctt	tgacacattc	tatatgagtg	ggtttactgg	taaattatgt	600
tattttttac	aactaatttt	gtactctcag	aatgtttgtc	atatgcttct	tgcaatgcat	660
attttttaat	ctcaaacggtt	tcaataaaac	catttttcag	atataaagag	aattacttca	720
rattgagtaa	ttcagaaaaa	ctcaagattt	aagttaaaaa	gtgggttgga	cttggaaca	780
ggactttata	cct					793

<210> 76

<211> 461

<212> DNA

<213> Homo sapien

<400> 76

accttgact	attccctca	gtccatctat	cgaggtcttt	gcaggaagca	tactgggaat	60
tgaaacgaga	gcctaaatga	catctaagaa	aggcagtggt	caataaccagg	tattaggtga	120
ggatgggatt	ctaaggacat	cagtgggagg	cagggagcca	ccttcagacc	tcagcatgga	180
agcttccaag	atccagagga	agaggcaaca	gcactgagag	tcataggtag	aagaatcatc	240
acagccctgc	taaccaggca	gctgatgccc	ctctccctcg	gtccctgtg	tccaaatcct	300
acaggggcat	ctgttggtg	aactcaacct	gaagccaaag	agaagatgag	tggagagagg	360
caacatttat	agagctcagg	tttctagggc	tggagaggga	tctggaggga	cacacaggag	420
acacctggca	taaccaaaaa	atgattaaaa	aaaaaaaaaa	a		461

<210> 77

<211> 642

<212> DNA

<213> Homo sapien

<400> 77

ggttgacga	aacacactgg	ggaatggagc	aaaacagtct	ttgaatatcg	aacacgcaag	60
gctgtgagac	tacctattgt	agatattgca	ccctatgaca	ttggtgggtcc	tgatcaagaa	120
tttgggtgtg	acgttggccc	tgtttgcttt	ttataaaacca	aactctatct	gaaatcccaa	180
caaaaaaaat	ttaactccat	atgtgttcct	cttgttctaa	tcttgtaaac	cagtgtcaagt	240
gaccgacaaa	attccagtta	tttatttcca	aatgttttgg	aaacagtata	atttgacaaa	300
gaaaaatgat	acttctcttt	ttttgctgtt	ccaccaata	caattcaaat	gctttttgtt	360
ttattttttt	accaattcca	atttcaaaat	gtctcaatgg	tgctataata	aataaacttc	420
aacactcttt	atgataacaa	aaaaaarawa	wattctttga	atcctagccc	atctgcagag	480
caatgactgt	gctcaccagt	aaaagataac	ctttctttct	gaaatagtca	aatacgaaat	540
tagaaaagcc	ctccctattt	taactacctc	aactggtcag	aaacacagat	tgtattctat	600
gagtccaga	agatgaaaaa	aattttatac	gttgataaaa	ct		642

<210> 78

<211> 519

<212> DNA

<213> Homo sapien

<400> 78
gcagaagaag aagcggacct tccgcaagtt cacctaccgc ggcggtggacc tcgaccagct 60
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gaaccggggc ctgcggcgga agcagcactc cctgctgaag cgcttgcgca aggccaaagaa 180
ggaggcgccg cccatggaga agcgggaagt ggtgaagacg cacctgcggg acatgatcat 240
cctacccgag atggtgggca gcatggtggg cgtctacaac ggcaagacct tcaaccaggt 300
ggagatcaag cccgagatga tcggccacta cctgggcgag ttctccatca cctacaagcc 360
cgtaaagcat ggccggcccg gcatcggggc caccactcc tcccgttca tccctctcaa 420
gtaatggctc agctaataaa aggcgcacat gactccaaaa aaaaaaaaaa aagggcggcc 480
gccaccgcgg gggagctcca cttttgttcc ctttaatga 519

<210> 79
<211> 526
<212> DNA
<213> Homo sapien

<400> 79
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ggtcacagcc tgatctctta tgtgttcata gccattoctt ctcccatcag aactgtttgt 120
cctgaatgtg ttctctagt tctagaaaat gaccactaat ttaaaaaact cggttgtgag 180
gtttgccagc aggcacttgt tccagaattt cccctcctgc ttcagccatg tccttgtcac 240
ttggcattct aagctaaaagc tttagcttcc caattogtga tgtgctaggc caagattcgg 300
gagctgttgc cagcctcgtc aaatatggaa gagaaacaac ctgcgggtcaa aaggagtgta 360
tttgtttaagt ggtgcgcgtc tatctcataa ctagatgtac caaccaggga agggccaagg 420
atggaaaggg gtaacttttg tgcttccaaa gtagctaagc agaagtgggg gagcagttta 480
gccagatgat ctttgattag gcaaacattg agttttaaag aggctg 526

<210> 80
<211> 281
<212> DNA
<213> Homo sapien

<400> 80
gttatattag tgggtagtgt aacattttat ccaggttggg gtgaggggag atggccacag 60
tagcaagtgg tgacactaaa taccattttg aaggctgatg tgtatataca tcattactgt 120
ccgtagcaat gaaggataca gtactgtgtt gtgggtgagt gttgctattg cccagcatta 180
atatttgggt gtgtatgttt gaggtatga aacacgcagg agtggttttt tgctattaat 240
tttaagagaa agcagctttt tcttaaaatt cactggttag a 281

<210> 81
<211> 405
<212> DNA
<213> Homo sapien

<220>
<221> misc_feature
<222> (1)...(405)
<223> n = A,T,C or G

<400> 81
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tagcaaaccg agcgatcatg tcgcacaaac aaatttacta ttcggacaaa tacgacsacg 120
aggagtttga statcgacat gtcattgctg ccaaggacat akccaasctg gtccctaaaa 180
cccatctgat gtctgaatct gaatggagga atcttggcng ttcagmagan tcagggatgg 240

<211> 401
 <212> DNA
 <213> Homo sapien

<400> 85
 cagtgtggtg gaattcccaa gatagaaatg aaaaactctt ttatagagtg ctgacatctg 60
 acattgagaa attcatgcct attgtttata ctcccactgt gggctctggct tgccaacaat 120
 atagtttggt gtttcggaag ccaagagggtc tctttattac tatccacgat cgagggcata 180
 ttgcttcagt tctcaatgca tggccagaag atgtcatcaa ggccattgtg gtgactgatg 240
 gagagcgtat tcttggttg ggagaccttg gctgtaatgg aatgggcata cctgtgggta 300
 aattggctct atatacagct tgcggaggga tgaatcctca agaattgtctg cctgtcatte 360
 tggatgtggg aaccgaaaat gaggagttac ttaaagatcc a 401

<210> 86
 <211> 547
 <212> DNA
 <213> Homo sapien

<400> 86
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 aaaacaaatc ctgtaactac ccagccagca agtatatagc acagaacact gtgttacttt 180
 acaagggctt atgtgactgg aataaggtgg tcccacttga ctgttcctaaa gagcagcttc 240
 tcagatcttc agtggtcact ggtaaatttc taacagtgtg tttgtgtaaa gtttgtcatt 300
 tcatactcca tacactacag ttgctgtcac tgatccctgt tttgctggct ttttaagctac 360
 ttgggtcaaaa atcctgcttc cttaaaacat agagaattaa tgagcatctc aagctttttc 420
 ttttcctttt taatgatgac tgcactatca agagtattct agtggttctc ctttgttttg 480
 catataatca tgcaccaaac tttttatttc ttttaaggtgg gagtatattt ttatttccta 540
 aatgcc 547

<210> 87
 <211> 530
 <212> DNA
 <213> Homo sapien

<400> 87
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 tttggcacct atgcgcctgc agaggttcct aaaagtaaaag ctctagaggc cgtcaaattg 120
 gcaatagaag ccgggttcca ccatattgat tctgcacatg tttacaataa tgaggagcag 180
 gttggactgg ccatccgaag caagattgca gatggcagtg tgaagagaga agacatattc 240
 tacacttcaa agctttggag caattcccat cgaccagagt tggtcgacc agccttggaa 300
 aggtcactga aaaatcttca attggactat gttgacctct atcttattca tttccagtgc 360
 tctgtaaaagc caggtgagga agtgatcca aaagatgaaa atggaaaaat actatttgac 420
 acagtggatc tctgtgccac rtgggaggcc atggagaagt gtaaagatgc aggattggcc 480
 aagtcacatc ggggtgtccaa cttcaaccac aggtctgctgg agatgatcct 530

<210> 88
 <211> 529
 <212> DNA
 <213> Homo sapien

<400> 88
 acctgagcta agaaggataa ttgtcttttg gtaactaggt ctacaggttt acatttttct 60
 gtgttacact caaggataaa ggcaaaatca attttgaat ttgttttagaa gccagagttt 120

atctttttcta	taagttttaca	gccttttttct	tatatataca	gttattgcca	cctttgtgaa	180
catggcaagg	gacttttttta	caattttttat	tttatttttct	agtaccagcc	taggaattcg	240
gtagtacttc	atgtgtattc	actgtcactt	tttctcatgt	tctaattata	aatgaccaaa	300
atcaagattg	ctcaaaagg	taaatgatag	ccacagtatt	gctccctaaa	atatgcataa	360
agtagaaatt	cactgccttc	ccctcctgtc	catgaccttg	ggcacaggga	agttctgggtg	420
tcatagatat	cccgttttgt	gaggtagagc	tgtgcattaa	acttgcacat	gactggaacg	480
aagtatgagt	gcaactcaaa	tgtgttgaag	atactgcagt	catttttgt		529

<210> 89

<211> 547

<212> DNA

<213> Homo sapien

<400> 89

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cacacaaggt	tatgattttt	ttaattactg	gcttctgatt	tctttcactt	ctgacccctt	120
tcctttttct	cagatgtagc	tgagtcttga	tcattttaag	acaacgatgg	gtagaatttt	180
gagattaatg	ttaattttcc	ctttttgtta	atttcagtc	cctctcacta	tgcttttgtc	240
cagaaggatc	aagaattcta	ccatcccttg	ggctcttgtg	tataaacaat	gttaaataaa	300
ggtagactca	gtctttaaga	tattagacag	tttttttagt	ccatgggatt	gtaaatataa	360
acattaactt	tcctataaga	atatatttggc	tttgtaatct	atagcctcaa	attggatttt	420
attatggatt	cactagacaa	acagctgttt	ccttattgtc	ttttttcttt	agtgtttctg	480
atttgctatc	agtagctgtt	tttaaagcca	tccaaggaaa	ataattattt	acagtttttg	540
aagtcac						547

<210> 90

<211> 528

<212> DNA

<213> Homo sapien

<400> 90

gagcagcaga	agctgtacag	caagatgatc	gtggggaacc	acaaggacag	gagccgctcc	60
tgagcctgcc	tccagctggc	tggggccacc	gtgcgggggtg	ccaacgggct	cagagctgga	120
gttgccgccc	ccgccccac	tgctgtgtcc	tttccagact	ccagggctcc	ccgggctgct	180
ctggatccca	ggactccggc	tttcgcccag	ccgcagcggg	atccctgtgc	acccggcgca	240
gcctaccctt	gggtgtctaa	acggatgctg	ctgggtgttg	cgacccagga	cgagatgcct	300
tgctttcttt	acaataagtt	gttgaggaa	tgccattaaa	gtgaactccc	cacctttgca	360
cgctgtgcgg	gctgagtgg	tggggagatg	tggccatgg	cttgtgctag	agatggcggt	420
acaagagtct	gttatgcaag	cccgtgtgcc	agggatgtgc	tgggggcggc	caccgcctct	480
ccaggaaagg	cacagctgag	gcactgtggc	tggcttcggc	ctcaacat		528

<210> 91

<211> 547

<212> DNA

<213> Homo sapien

<400> 91

atataaccatt	taatacatatt	acacttttctt	atttaagaag	atattgaatg	caaaataatt	60
gacatataga	actttacaaa	catatgtcca	aggactctaa	attgagactc	ttccacatgt	120
acaatctcat	catcctgaag	cctataatga	agaaaaagat	ctagaaactg	agttgtggag	180
ctgactctaa	tcaaatgtga	tgatttgaat	taraccmttt	ggscyttgra	ccttymtwrg	240
raaaawgrmc	cmaccttttyt	taacmtgrac	cwccytmatc	tctagaagct	gggatggact	300
tactatyctk	gttwatattt	taaatackga	aagggtgctat	gcttctgtta	ttattccaag	360
actggagata	ggcagggcta	aaaagggtatt	attatttttc	ctttaatgat	ggtgctaaaa	420

```
<210> 92
<211> 527
<212> DNA
<213> Homo sapien
```

```
<220>
<221> misc_feature
<222> (1)...(527)
<223> n = A,T,C or G
```

```
<210> 93
<211> 531
<212> DNA
<213> Homo sapien
```

```
<210> 94
<211> 547
<212> DNA
<213> Homo sapien
```

```
<220>
<221> misc_feature
<222> (1)...(547)
<223> n = A,T,C or G
```

<400> 94
gttaaacatg gtctgcgtgc cttaagagag acgcttcctg cagaacagga cctgactaca 60
aagaatgttt ccattggaat tgttggtaaa gacttgaggt ttacaatcta tgatgatgat 120

```
<210> 95
<211> 1265
<212> DNA
<213> Homo sapien
```

```
<210> 96
<211> 568
<212> DNA
<213> Homo sapien
```

<400> 96						
ccagtgtggt	ggaattcggt	ttaattacaa	aatttgatca	cgatcatatt	gtagtctctc	60
aaagtgtctc	agaaattgtc	agtggtttac	atgaagtggc	catgggtgtc	tggagcaccc	120
tgaaactgta	tcaaagttgt	acatatttcc	aaacattttt	aaaatgaaaa	ggcactctcg	180
tgttctctct	actctgtgca	ctttgctgtt	ggtgtgacaa	ggcatttaaa	gatgtttctg	240
gcattttctt	tttattttgt	aggtgggtgg	aactatggtt	attggctaga	aatcctgagt	300
tttcaactgt	atatatctat	agtttgtaaa	aagaacaaaa	caaccgagac	aaacccttga	360
tgctccttgc	tcggcgttga	ggctgtgggg	aagatgcctt	ttgggagagg	ctgtagctca	420
gggcgtgcac	tgtgaggtcg	gacctgttga	ctctgcaggg	ggcatccatt	tagcttcagg	480
tgtcttgcct	tctgtatata	gtgacatagc	attctgctgc	catcttagct	gtggacaaaag	540
gggggtcagc	tggcatgaga	atatTTTT				568

<400> 97

<210> 98

<211> 547

<212> DNA

<213> Homo sapien

<400> 98

tactgggtgc	caagctatgt	gccaggcact	ttacatgtat	tgattttaaca	cttaacagcc	60
actctatatt	attccctttt	tacagatgag	gcaatttaag	ctcaaagcat	ttaagtagac	120
aaccaaccta	gaatcacata	gcaaatgaca	gaagccagag	gcctcccaag	tctctctaac	180
tccaaaccct	atgcttactc	tactatatca	cactaccttg	caataggaca	aagggaatat	240
gtggtaaaact	atgttcccag	catctaaaag	ccaggagtgg	ttttcatttt	tctttaagaa	300
gatgatagtg	tgatttgaaa	catatctgaa	tttcagaaga	ggggactttt	aaaaattgcc	360
actcataagg	aaagaaagaa	ctttttcaca	tatttttgaa	agaaacgatg	gtgagaagat	420
attcttgata	atagagatat	gctaacattt	gctttgggtg	ttttgtaggt	tagatttttt	480
tgggtgtgtac	tttataggct	tgcatattgc	ttacttttaa	cagctgaagt	tctaagtaag	540
agtgttc						547

<210> 99

<211> 122

<212> DNA

<213> Homo sapien

<400> 99

```

cagcctttct gtcacatct ccacagccca cccatcccct gagcacacta accacctcat      60
gcaggcccca cctgcccaata gtaataaagc aatgtcactt ttttaaaaca aaaaaaaaaa      120
aa                                                    122

```

$\langle 210 \rangle$ 100

<211> 449

<212> DNA

<213> Homo sapien

<400> 100

ctgacggcctt	tgctgtccca	gagcgccta	aacgcaagaa	aagtcgatgg	gacagttaga	60
ggggatgtgc	taaagcgtga	aatcagttgt	ccttaatttt	tagaaagatt	ttggtaacta	120
ggtgtctcag	ggctggggtt	gggtccaaag	tgtaaggacc	cctgcctt	agtggagagc	180

tggagcttgg	agacattacc	cottcatcag	aaggaatfff	cggatgtfff	cttgggaagc	240
tgffffggtc	cttgggaagca	gtgagagctg	ggaagcttct	tttggctcta	ggtgagttgt	300
catgcgggta	agttgaggtt	atcttgggat	aaagggctct	ctagggcaca	aaactcactc	360
taggtttata	ttgtatgtag	cttatatfff	ttactaaggt	gtcaccttat	aagcatctat	420
aaattgagtt	ctffffctta	gttgtatgg				449

<210> 101
 <211> 131
 <212> DNA
 <213> Homo sapien

<400> 101	
ccatgtttctc	tcttgaactac
gcataatgtga	gatttgcccc
tccgccccgc	tcgtgatagc
catccagatc	ttttacctgg
ccctgtcttg	gagaatctgt
tttcaatctc	cactgattgc
ccccttgctg	g
	60
	120
	131

<210> 102
 <211> 199
 <212> DNA
 <213> Homo sapien

<400> 102	
ctgctgcgcc	tgatgctggg
acagccccgc	tcccagatgt
aaagaacgcg	acttccacaa
acctggattt	tttatgtaca
accctgaccg	tgaccgtttg
ctatatctct	ttttctatga
aataatgtga	atgataataa
aacagctttg	acttgaaaaa
aaaaaaaaaa	aaaaaaaaaa
aaaaaaaaaa	aaaaaaaaaa
	60
	120
	180
	199

<210> 103
 <211> 321
 <212> DNA
 <213> Homo sapien

<400> 103	
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tttatttgca	tattaaaaaa
attgtgcatt	ccaataatta
aaatcatttg	aacaaaaaaa
aatggcactc	tgattaaact
gcattacagc	ctgcaggaca
ccttgggcca	gcttggtttt
actctagatt	tactgtcgt
cccaccccc	cttctttcac
cccacttttt	ccttcaccaa
catgcaaagt	ctttccttcc
ctgccacca	gataatatag
acagatggga	aaggcaggcg
cggccttcgt	tgtcagtagt
tctttgatgt	gaaaggggca
gcacagtcac	ttaaacttga
t	
	60
	120
	180
	240
	300
	321

<210> 104
 <211> 309
 <212> DNA
 <213> Homo sapien

<400> 104	
tttttttttt	tttttatfff
tttttttgca	tcaaaaaact
ttatttccat	ttggcccaag
gcttggttagg	atagttaaaa
aagctgccta	ttggctggag
ggagaggctt	aggcaaaacc
cctattactt	tgcaaggggc
ccttcaaaag	tctctgggct
tctatttcaa	cgcgatgat
gtggctctgg	aaggcgtgag
ccactttttc	cgggaactgg
ccaaggaaaa	gcccaggggc
tacaaccgtt	tcctgaaaat
gcaaaaccag	cggggcggcc
gcgctctttt	ccaggacatc
aaaaagcca	
	60
	120
	180
	240
	300
	309

<210> 105

<211> 591
 <212> DNA
 <213> Homo sapien

<400> 105
 cttattttctg catggggtcgg agagtgggag ggactgcttt actgagttat agtgaatgta 60
 gttttaacct aagcgctca catgactaac tctcatcca tcaagaatga gctcagctct 120
 cacttcccca ctctcacc cctgtaaag taacctttct ccaaggttat gcttcaacag 180
 gaatagctaa catttattaa attgtggcac gtaagtatct tggatatatt ggctcattga 240
 atcctcacac ctactatttt acagagatgc cagtggggct tgagattgaa tcacttgccc 300
 aggcctccac tgctggtaaa cagtagaggg ggctcctgac ccatcagtct ggcttgacaa 360
 cccattccct caactgcgga tcccggaatc ccttatcacc ctgttgattt ctccataggc 420
 tgtggttaaca tttgttgcac gaatggaccg ttgaaatagg gcttggcagg gagaaattca 480
 ggaaatgaat gaatggttct tccctggcag cctttgatga cttacaagcc ccttcaaggg 540
 ggaaagccat ttttctccct gggactcctt gaaagcccgg gagccctgcc t 591

<210> 106
 <211> 450
 <212> DNA
 <213> Homo sapien

<400> 106
 ctgccactcc tgctctgct accccgaaac cggagaggga gctcaataat aacacaggtc 60
 ccactaaact aattaagggt ttggcataac ctgtcattga attcaagtgt ccaacaactg 120
 tttgcttaaa atatcattag acctaataatt tttttcaaag gcacaaagtt taaacatggg 180
 gggggcgggt gttgagaggg gtctgggata cccttaaacc caaaaaagtg atttgttccc 240
 ccttgcccag aagggtgact gttccactgg gctgtcacc acaggacatt ttccatgaca 300
 agcactcacc ttcttgggga aggggcatca gggtggcaca ggaaaggccc aagtgagggg 360
 ccactctgta cattaatact ttggtgatta atgtttgggg agaggcagga ttctcaccca 420
 cctttttgac ttcaaacact ctcaactcaag 450

<210> 107
 <211> 116
 <212> DNA
 <213> Homo sapien

<400> 107
 tcgacgaaag ttactgtcac tcagttgtaa atccatcagc ttttcacctg ttaaaaattt 60
 tgcaaaatat acatgttctc ctctgtttt caattcttcc atcttttttc ttgagg 116

<210> 108
 <211> 291
 <212> DNA
 <213> Homo sapien

<400> 108
 ctgctogaag ttgtcaaaac ccacgtgcag ggcaatggag agtccgatgg ccgaccacag 60
 cgagtagcgt cctcccacc aatcccagaa ctgcaacatg ttttgagggt caattccaaa 120
 ctcccttcaact ttggttgtgt tagtagacag ggcaacaaag tgcttcgcca ctgcagtagg 180
 atccttggcc gcttgagaa accactcctt cgccgtctct gcattcgtga tggctctctg 240
 ggtagtaaag gtcttggagg caatgatgaa cagggaggac tcgggggttca g 291

<210> 109
 <211> 662

<212> DNA

<213> Homo sapien

<400> 109

gctgtttcca	cagtacgcct	gcctcacacc	ttgcgatgcg	ccaacatcac	catcattgag	60
caccagaagt	gtgagaacgc	ctaccccggc	aacatcacag	acaccatggg	gtgtgccagc	120
gtgcaggaag	ggggcaagga	ctcctgccag	gggtgactccg	ggggccctct	ggctctgtaac	180
cagtctcttc	aaggcattat	ctcctggggc	caggatccgt	gtgcgatcac	ccgaaagcct	240
ggtgtctaca	cgaaagtctg	caaatatgtg	gactggatcc	aggagacgat	gaagaacaat	300
tagactggac	cccccacca	cagcccatca	ccctccattt	ccacttggtg	tttggttcct	360
gttcactctg	ttaataagaa	accctaagcc	aagaccctct	acgaacattc	tttgggcctc	420
ctggactaca	ggagatgctg	tcacttaata	atcaacctgg	ggttcgaaat	cagtgaagacc	480
tggattcaaa	ttctgccttg	aaatatgtg	actctgggaa	tgacaacacc	tggtttggtc	540
tctgttgat	ccccagcccc	aaaagacagc	tcctggacct	tgccccgggg	cggcccgcctc	600
ggaaaggggg	cgaaatttct	tcaagaatat	ttccatttcc	acaaacttgg	ggccgggggc	660
cc						662

<210> 110

<211> 323

<212> DNA

<213> Homo sapien

<400> 110

tcctgtgaaa	cagcccattt	tcctacctac	tgtgggttgc	tgctcaggag	gaacgatata	60
cgccaatata	agcaggaaat	ctgcagctcc	tctgctatgt	gcctcagaac	actttcaatt	120
tttctgggtca	atgctctgat	taggtatcat	acataaaaagc	cagcatatta	gtttaaatct	180
ctaacaaaaa	actatatttt	ccaaagtcat	tatcatttgg	gccaatataag	tgatcttttc	240
gtgctttgtt	gagcttcctc	tttagggcat	ctcttctttc	ttcccattca	tgaagttcgg	300
catttccatg	tgcaaattta	cag				323

<210> 111

<211> 336

<212> DNA

<213> Homo sapien

<400> 111

tccagtgcgc	tccagcctta	tctaggaaaag	gaggagtggg	tgtagccgtg	cagcaagatt	60
ggggccctcc	ccatcccagc	ttctccacca	tcccagcaag	tcaggatata	agacagtcct	120
cccctgaccc	tcccccttgt	agatatcaat	tcctaaacag	agccaaatac	tctatatcta	180
tagtcacagc	cctgtacagc	atttttcata	agttatatag	taaatggctc	gcatgatttg	240
tgtctctagt	gctctcattt	ggaaatgagg	caggcttctt	ctatgaaatg	taaagaaaga	300
aaccactttg	tatatatttg	aataccacct	ctgtgg			336

<210> 112

<211> 218

<212> DNA

<213> Homo sapien

<400> 112

tttttttttt	tttttttttt	tccagtcagg	agtattttta	atcactgtct	acagagacac	60
ctacatacac	acacgggtgg	ggaatgaacc	caaagttttt	aggatgaagtc	tctcagggcc	120
caccccgctg	cacagacctt	cctcggttgc	agagattctg	ggcaaagcat	ccgtgctctc	180
atgagattat	cctggggaga	tttagaagaa	ttttgtgg			218

<210> 113
 <211> 533
 <212> DNA
 <213> Homo sapien

<400> 113
 ctgcaccgac agttgcgatg aaagttctaa tctcttccct cctcctgttg ctgccactaa 60
 tgctgatgtc catggtctct agcagcctga atccaggggt cgccagaggc cacagggacc 120
 gaggccaggc ttctaggaga tggctccaga aaggcggcca agaattgtgag tgcaaagatt 180
 ggttcctgag agccccgaga agaaaattca tgacagtgtc tgggctgcca aagaagcagt 240
 gcccctgtga tcatttcaag ggcaatgtga agaaaacaag acaccaaaagg caccacagaa 300
 agccaaacaa gcatcccaga gcctgccagc aattttctcaa acaatgtcag ctaagaagct 360
 ttgctctgcc tttgtaggag ctctgagcgc ccactcttcc aattaaacat tctcagccaa 420
 gaagacagtg agcacaccta ccagacactc ttcttctccc acctcactct cccactgtac 480
 ccacccttaa atcattccag tgctctcaaa aagcatgttt ttcaagatct aaa 533

<210> 114
 <211> 261
 <212> DNA
 <213> Homo sapien

<220>
 <221> misc_feature
 <222> (1)...(261)
 <223> n = A,T,C or G

<400> 114
 ccatatctgc tcggcgctac ttctttcttg gattgatcct gantgatgca ttggcgatgc 60
 ctttgagaaa ggacatgtga tgtgatggtc ttcacgttcc acatgtactc gggcaaatag 120
 ggggacaaac tgaagttaaa caggtcgaaa ctagaggagc tgctgaccct ggagctgacc 180
 actttcttgg ggaaaaggac acatgaaggt gctttgcaaa agctgatgag caatctggac 240
 accaaccatg gacaacaacg t 261

<210> 115
 <211> 267
 <212> DNA
 <213> Homo sapien

<400> 115
 cctctcctgt gggttccaga ccctgttcca gcaacaattg ctgggacacc tgggccgact 60
 gctccacctc gccaggccct ggccctctcc atctcagccc tgacagccac ccagtataa 120
 acacagcagg cttcctaagc aatgtgacgc accagagggg tgggtggtaca cgttcccctt 180
 gaagtcactt gaaaattaga gaacagattt gcctcatagc tgaagagaga ccctattcca 240
 agcatgaatg gccttgacaa tgttcct 267

<210> 116
 <211> 239
 <212> DNA
 <213> Homo sapien

<400> 116
 ctgatgacct ggggtctagt gaaaatgcag ggtcagattc agtgggtctg gggctctgaat 60
 ctctaaggcg ctgccaaagt atgctgatgc tcctggcttg tggaccacc cgtgtatagc 120
 aaagctctag actaggaggt ctcaaccttg gctgcacaga attatctggg gagtttttaa 180

atttcccagt gccagggctg cattcatatc atagtagaga cagggttttg ccatgctgg 239

<210> 117
<211> 168
<212> DNA
<213> Homo sapien

<400> 117
aaaaaacttt tatattgctg catcttccac agttctttgg gtagtctctg aacttaaaat 60
ttgtaggagt ttagactac ctaaattttt aagttatgga tttgttcata ggtttaggg 120
gtaggtaaaag aaggaaacag acaagaaaat ggcttcttga ggtggcag 168

<210> 118
<211> 150
<212> DNA
<213> Homo sapien

<400> 118
aaaaaaaaaga gtttatttag aaagtatcat agtgtaaaca aacaaattgt accactttga 60
ttttcttggga atacaagact cgtgatgcaa agctgaagtg tgtgtacaag actcttgaca 120
gttgtgcttc tctaggaggt tgggtttttt 150

<210> 119
<211> 154
<212> DNA
<213> Homo sapien

<400> 119
aaactgtgtg agatattaac cagccgccct gttataaaat caggaaatcc aaacagcgat 60
ttacaccgat taacaccccc ttttatattt tttcaaatac actgagaaaa taatcaaacg 120
ttttcatctc tcttgtcttt tttgtttttt tcct 154

<210> 120
<211> 314
<212> DNA
<213> Homo sapien

<400> 120
ctgctgtggag tgacgggagg agggaatcac tgtgtgtgcg agagtgttc agactcaatt 60
tccaaaataa ttttcacccc tctaagcatg taaattcaaa gatggatcct tcatagaaat 120
taaaaaatca atttgagctc atttgaata cagaacaagt atggcacaga tggaagtcct 180
gccacgtttc ctttaatgat gctgactctt gtatcacaca ggccagcatg aagtttctta 240
ctcagacttt acaggcattt tccgtaattc aatcagtcct gctcccagca caacacagga 300
ggtgattcga gaat 314

<210> 121
<211> 601
<212> DNA
<213> Homo sapien

<400> 121
aaaaaaaaacc taattcattg aagtaataac caaataattt tcaatcttga ttcaactgtg 60
attcaaactc tacaccattt gcccttctta tgaatttatg tataaaattt ttttaagagtc 120
agagtttttt tttcttgatt aattggatgt atttcacaga atttccaact gctcacgtta 180

```
<210> 122
<211> 486
<212> DNA
<213> Homo sapien
```

```
<210> 123
<211> 239
<212> DNA
<213> Homo sapien
```

```
<210> 124
<211> 610
<212> DNA
<213> Homo sapien
```

<400> 124						
ccanccaagt	cnttgatgat	cactgaccen	cgcgcgctg	ctggaccaag	gtggctgcgg	60
ggaaatcgcc	acngngcttt	cggttttctt	ggtgaaggaa	tacaccgcgc	cgacagcagg	120
ttttcagtc	gggtcaggg	ctgttgcttg	cgcgcgaaaa	tcaccggtac	gccgaggttc	180
aggccgggtc	tgatcgccgg	tgcaatgcc	gaggcttcga	tggtgacgat	cttgggtgatg	240
cccgaatcct	tgaacaacgc	agcgaattca	tcaccgatca	gtttcatcag	cgccgggtcg	300
atctggtggt	tcagaaaggc	gtcgaccttg	agtacctgat	cggaaagcac	gatgccttct	360
tcgcgaattt	tcttqtgcag	tgcttcacg	aaagcttcct	ctgttggcgc	aacacgcgcc	420

gaaagtagat taaaaagtag tcgattctag cgctttaaca tcgcgcgtat atccgccagg 480
 gcggtattgc cgcgaacggc tttgacttcg gttggtgtgt cgtcggtgcc ttcccatgcc 540
 aggtcatccg gcggcagttc gtcaaggaac cggctggggg cacaatcaat gatctcgccg 600
 tactgcttgc 610

<210> 125

<211> 196

<212> DNA

<213> Homo sapien

<400> 125

ctatagggct cgagcggccg cccgggcagg taaaaaatca gcccctaatt tctccatgtt 60
 tacacttcaa tctgcaggct tcttaaagtg acagtatcct taacctgccca ccagtgtcca 120
 cctccggcc cccgtcttgc aaaaagggga ggagaattag ccaaactctg taagctttta 180
 agaagaacaa agtttt 196

<210> 126

<211> 247

<212> DNA

<213> Homo sapien

<400> 126

aaattagtta aaaaaatgca ttcctcattt gatatagcca cattccaaat gcttaaaagc 60
 cgcatgtatc tagtgactac catactggag agtacaaata tagaacttta cccgtcactg 120
 cagacagttc tggttgattg tgcagcattg gacaatatat acagtttgcc tgtatatgag 180
 aaagagagag agagagagag tgtgtgtgtg tgtgtgtgtg tgaagtgcaa taaggctgac 240
 aggcac 247

<210> 127

<211> 590

<212> DNA

<213> Homo sapien

<400> 127

cctccacggc atggcgcaat tgttgttcag gggccgccag gttgctgcc atgccgatgt 60
 agatacgttc cactgtgctta ctgccagac gcactcgaag cgtcgccagc gctacgtttg 120
 cgcttgctgc cactgctgcg gcgacgctt ttcgggccat cgccggtggc ttgcgcttg 180
 ctgctgagct ctttgatcat ctgcggcgc tggctgtcgt tggcgtcctg gtagtcggtc 240
 caccactcgc caaggccgtc ggtctgttcg ccggcgcttt cacgcagcag caggaagtca 300
 tagcccgga cgggaagcgcg ggttgctccag caacaggtcg gcacgtttgc cgctgcggcg 360
 tggcaggcgc tcctgcatgt cccagatttc acggatcggc atggtgaagc gtttcgggat 420
 ggcgatgcgc tggcattgct cggcgatcag ctgctgagca gcttcctgca tggctggaat 480
 tgccggcatg ccacggctct gcaggcgcat gacgcgtttc gaaagcgcgg gccacaacag 540
 ggccggcaaag aggaacgccg gggtgaccgg tttgttctgc ttgatgcgca 590

<210> 128

<211> 361

<212> DNA

<213> Homo sapien

<400> 128

ctgcccacgg aaaccctcca ggagctgctg gacctgcaca ggaccagtga gagggaggcc 60
 attgaagtct tcatgaaaaa ctctttcaag gatgtaacca aagtttccag aaagaattgg 120
 agactctact agatgcaaaa cagaatgaca tttgtaaacy gaacctggaa gcatcctcgg 180

```
<210> 129
<211> 546
<212> DNA
<213> Homo sapien
```

```
<210> 130
<211> 733
<212> DNA
<213> Homo sapien
```

```
<220>
<221> misc_feature
<222> (1)...(733)
<223> n = A,T,C or G
```

```
<210> 131
<211> 305
<212> DNA
<213> Homo sapien
```

```
<220>
<221> misc feature
```

<223> n = A, T, C or G

aaacacatac	gaatanttna	actgtgatta	tgaagtgaca	gccggctaaa	tatgtcttgt	60
attttctctc	ttcctttttt	tgctaactca	tcctttattc	cattcctgct	tcocatggtaa	120
tgcaggctca	aataaattac	taggatacaa	gattacttca	agcctctttt	ctgttggaact	180
cataatatga	taagcatttg	ttacaagatt	gcctgtagtt	gttttagggga	caaattatat	240
tagggaaaaga	aagtctttct	ttagttgggt	aaattttcta	ttataattgg	gtactaaatt	300
tatttt						305

<213> Homo sapien

aaacaatgct	acactcattt	ttggcaaaagt	gctgtattgt	tcagtcctgtg	tacaaaaactg	60
accatctatg	aaccaatcag	tataaaaaaat	ttctataaaa	acaaaattta	gacagcggct	120
caagaaaaca	agctgccatt	tatgcataga	ttgatgtaca	gtaacctaac	caaatgtccc	180
ttttgaattt	tcaagttact	gaaaaaaaaat	gtgtcgagaa	acacattaag	aaggcacatg	240
tacagctctac	aatactcttc	agtcctcccta	actcatgccc	tgccctata	aaggaaatat	300
gttcacaatt	ttacttgaga	aaaaaaaaaca	aagccactta	aaaaaaaaaaa	aacacacacg	360
caattattaa	agttcaaaat	ctctggagga	aaatacaagc	aaaaccactc	atacactcca	420
agcctgaaac	acacatctaa	cctcccagg	tactggtttg	gttttcagag	gtccacctag	480
aaaacaaatc	taaaacttca	ggcaaaacag	agcaaaactg	gacattttaac	aattacacaa	540
ttttt						545

<213> Homo sapien

<223> n = A, T, C or G

aatatattatt	actaatatct	tataatgttt	tgtggnacca	tggcataacct	tgggtactat	60
tgtaacanat	agttcaggaa	accctactat	aagggtttatc	aaatgggtctc	ataaacagtt	120
acttattcaa	gcacgccaaa	gctcagtgaa	aagtatttttt	cacccttact	ctttctcgtg	180
tcattcaaag	agaagttttg	atgtagtgtg	tttattttgta	gggagtaatg	aacagatcca	240
tttcacagta	gacttttgtc	tctaggtgat	gcagctaatt	gccccagttt	ggaaaacatg	300
gacttqqatg	aattgtcttt	tgtttgggac				330

<213> Homo sapien

$\langle 222 \rangle$ (1) ... (627)

<223> n = A,T,C or G

<400> 134

```

aaatattact tcaaatacat tttaaagctc aacaaacttg tgttgaactg aattgcagat      60
cctgaactct atttgaaaat acatcatgaa acagaaaanc ccattccaaa tgaaaatgat      120
agtgccttgt tgggggtggg aatgaggcgg ggagactaaa tcactattaa cagacttctt      180
ttcccaatgc aatttgtcaa aagttcaaaa gttctgaaat gtactaaatc ttaagcaaatt      240
taaattcatg atattactaa aactttttta atagtgcatt gacttatcaa gttatagtgg      300
ctgcattaag aacaaattat tgtgtgaaat acctgtataa acacaaaata caattaaata      360
tttctttaca aaaagctgag cattacgcac aatagtggaa tgtctttcat taggtgtatt      420
ttttaaagat taacaaaagt aacatttcct aaaatgtata catgtgccat atttttgcaa      480
acatgcctga gaatgtattt aaaacatttc tgtagtaaga gtttgcaaga acttcacaaa      540
cctgcaataa aaatgcatct ttttaaaaag gtgaaaatgg catctccaca ctgcaacaat      600
tcaaaaagtg cagcatccct aatctttt                                     627

```

<210> 135

<211> 277

<212> DNA

<213> Homo sapien

<220>

<221> misc_feature

<222> (1)...(277)

<223> n = A,T,C or G

<400> 135

```

aaaatcaaatt atattatttg ttaaaaatca gcttgtttca ttacnggaaa ttacaccagt      60
ccgttctatt tactttcaaa ccatattcaa ctctcaact ttcaaactg taatcaacta      120
atttcaaaaag ggaaaaggta ccctttataa aggagagatc tgtaagaca ccaagaaatc      180
aaaattaata tcacttaata attaagtggg taacacatgc ctccaatac agtgcagtga      240
gaaacacaaa acatcaattc ccgcgtactc tgcgttg                                     277

```

<210> 136

<211> 486

<212> DNA

<213> Homo sapien

<400> 136

```

aaaacagaat gaattcattg ttacagttac agaagtcaga agcccaaata cagtctgcct      60
gaaccaaagc cagggtcagc aagggttcctt tccactgttt tgccaacttc tagaggccac      120
ctgtattcct tgggttcattg cccctctctt catcatcaaa taatcagcat agctttatga      180
cattggcagc tctgattttg ctcttttgcc ttcctcttat gtagaccctt gtaattacat      240
tgggtacacc cagataaccc caaataatct ccctatctca agattcttaa tgtaattata      300
ttgggaaagt cccttttgct atataagata acatagcaat ggattccaag gattagtatg      360
tgagtttctt ttgaggggct ataattaacc ctaccacaat atggaaatgt ctattgtttt      420
tctatgtacc agaaataaga cattaggatg tgaaattaat aacataacac cacttacggc      480
atcacc                                     486

```

<210> 137

<211> 552

<212> DNA

<213> Homo sapien

<220>

<221> misc_feature
 <222> (1)...(552)
 <223> n = A,T,C or G

<400> 137
 ccattcttgca tcaaattgttc ttaaggcagt gactggctat caaccacagt ttctgtctcc 60
 ccagttgcaa acacaggatc catgcaacag ttctgagacc atacacttag aaaccacagg 120
 ggatgcggat caaatgcaga actcccaaact tataaaacag tcagggtaca ctcaaaacaa 180
 aacatagaac atcaacaaca cacatctccc aaaaaagaag tgcaacgcat gcttgtataa 240
 accaacaata acaaaaaaac cacaataaaa aatgcagagt ctcccaaaca agttttcaaa 300
 tgtattgcan aaagaaaaaa aatgtatata tatataaaat taaaaagtct gaaatactag 360
 tgcatagtca attacctaac accaagtttc ttttctttct gtccaagctc tactgcccct 420
 ctgatactag cagcatgtct acaggctaag accatagcag caaaaaacgt ttttcatttg 480
 gcattttaca aattaaatta ctgaataaaa atataatttt ttataaaaact atttcttaca 540
 gtaataattt tt 552

<210> 138
 <211> 231
 <212> DNA
 <213> Homo sapien

<400> 138
 aaattttact agtgttactt aatgtatatt ctaaaaagag aatgcagtaa ctaatgcctt 60
 aaatgtttga tctctgtttg tcattacttt ttcaaaatat ttttttctgt aaagtataat 120
 atataaaact tcttgcttaa attgaatttc tatattagtg gttaattgca gtttattaaa 180
 gggatcatta tcagtaattt catagcaact gttctagtggt tttgtgtttt t 231

<210> 139
 <211> 535
 <212> DNA
 <213> Homo sapien

<400> 139
 cagttgccaa ccctctgaac cgtttaggcc ggttcacgc tgcccttgaa tctggggcgg 60
 tggatgatccg gcaaggggtg aaaccaaaga gcgggggctg tgaggccctt cgcagtcctt 120
 cgtaagtgcg tgcgatggag tgaactatca cgcacgtgt ttatttcgtc aacacgaaat 180
 gtgatttatt tttgcgaatt aacacggcag ttctcgggta cgttttcgga aagcgtggga 240
 tatgattctg tctatcctgt acggatatac agtaattacc gggaggggat tccatggcga 300
 agaagcaggc ggcaccggca gcacggcagg aaatgagcgg tatggcgcgc ctcgggcttc 360
 gcgtctcatc gatgattaat caccgggtcg cccagacgca gcgctgggtt acgattcatc 420
 gcctggacac ggatggggat cgggagtggt aagaggttct gagcgtgatc gctgataccg 480
 acgagctcga gctgacgctc aatgacgatg gcagtgtgac ggtgaggtgg gagca 535

<210> 140
 <211> 640
 <212> DNA
 <213> Homo sapien

<220>
 <221> misc_feature
 <222> (1)...(640)
 <223> n = A,T,C or G

<400> 140

```

acattggtgg cacttgaact gagtgcaaac cacaacattc ttcagattgt ggatgtgtgt      60
catgacgtag aaaaggatga aaaacttatt cgtctaattg aagagatcat gaggagaag      120
gagaataaaa ccattgtttt tgtggaaacc aaaagaagat gtgatgagct taccagaaaa      180
atgaggagag atgggtggcc tgccatgggt atccatgggt acaagagtca acaagagcgt      240
gactgggttc taaatgaatt caaacatgga aaagctccta ttctgattgc tacagatgtg      300
gcctccagag ggctagggtta gtacaaactc gcattcatgg cttgggtttcc cagaagatct      360
ccatttaact tttttaaaga aagtttattg ctttctttaa cctgcatttt ttctaagttt      420
tttttcgcat aaaggtgctg tctttgtggc aaggcctagg catgacaatc ggaggactcg      480
agggggatgg aggactagtg atccggctgg ctgcttccag tcgattagag aggtgaaaaa      540
gctgaacgtg tgcccantna atcttcaaaa aggcagaaac atatcacctt ntgccccnt      600
aaacttgttc tttttccgaa ggggaaaaaa aaaatggaaa      640

```

```

<210> 141
<211> 127
<212> DNA
<213> Homo sapien

```

```

<400> 141
aaaaatcaca cactgacaac acagaaatac gaaatgctag gaaaagtcta gcatatgaag      60
gaaaaacatg tcttatgcac tctaataataa ttttttcaat tagtataaag gcaaatgcgg      120
ttttttt      127

```

```

<210> 142
<211> 126
<212> DNA
<213> Homo sapien

```

```

<220>
<221> misc_feature
<222> (1) ... (126)
<223> n = A,T,C or G

```

```

<400> 142
aaatatactc tggatgcntt caagtaatac taatcatttc atgnngnaaaa gtcttttaat      60
aaacaaattc agagtaaaat taattgaaat atttataata catttggttac acagttattt      120
ccaata      126

```

```

<210> 143
<211> 730
<212> DNA
<213> Homo sapien

```

```

<220>
<221> misc_feature
<222> (1) ... (730)
<223> n = A,T,C or G

```

```

<400> 143
gcaagttctg gagggttcac ttctgagcct gaattccctc ccctgcaaaa tgggggaata      60
ccctcctcag aggggtccctg cgagggtgag gggagatcag catggcagggt gtgctgggca      120
cggcagggcc tgggaagggc agatcctttc cccatccctg ccacaaacaa cccaaacctt      180
taaaggagag caatggcctt gtgtcaaaaa caaaaacaaa acaaaaccct gtcctaggag      240
actggggccc taattttctaa tagcaagcct ttatgagtcc ctaacactct actgggctga      300
gtatctcaca cgccagagga taacctgcct tctgctcacc accaccccggt agtagttgtc      360

```

atttgtgtcca tttcacagat gaggcaaagg ctcagaagag tcatgtgtta aaccagcttc 420
tagagcccat gcaggagctg caggtgggga gaatcacctc taggtgctct tcccatggaa 480
tcctcaccct ccttgagtgg tcactcactc anctttccaa tgggtgtgtg acctttgacc 540
agctttcttt ccttntctgg gcctcagttt cccaccttgg acaaagtaag aggtctcttg 600
ggnttcangg tagttcttcc taacttcttt tccttttcat ttgagcatcc ttcttcattt 660
tttgccacct ctcttgtcat tacangcttt taccttcggc cgcgaaccac gcttaagggc 720
naaatttcca 730

<210> 144
<211> 485
<212> DNA
<213> Homo sapien

<400> 144
ctggtcagaa atgattctct tgtgacacca tcgccacaac aggcctcgggt ctgtcctccc 60
catatgttac ctgaagatgg agctaccttt cctctgtgtg gcattttgtc gcttatccag 120
tcttctactc gtagggcata ccagcagatc ttggatgtgc tggatgaaaa tcacctgtgt 180
tgctgtgtgg gtctgtgtcc gccacttcta atcctcatca tgacaacgtc aggtatggca 240
tttcaaata agatacaacc attgaaggaa cgtcagatga cctgactgtt gtagatgcag 300
cttcactaag acgacagata atcaaaactaa atagacgtct gcaacttctg gaagaggaga 360
acaaagaacg tgctaaaaga gaaatggtca tgtattcaat tactgtagct ttctggctgc 420
ttaatagctg gctctggttt cgccgctaga ggtaacatca gccctcaaaa atattgtctc 480
aacag 485

<210> 145
<211> 465
<212> DNA
<213> Homo sapien

<400> 145
ccaagacagc tcgtttctgg agagtatgag ggtgtgtttt cttattgtga aaggaactac 60
cttctcttag agggtaggaa gaatgtgggtg tgtgtgtgtc tcataaagca accggacatt 120
ataggtgccc aggtcatcta taaaaacgat ccttgggctg tgtaaaaatg aagtggcttt 180
tcagtatcct ctttcacact tgctgcttcg ggagactatg caatgatggg aaggtgattg 240
cccctttatt tcattcagtg ccattggtccc tgttgttgta gtaatttatt tgtttagttc 300
atTTTTTTTT tcttaacagt caaggggaag agtgattcct cactactgctt tcaagctgga 360
ctgagccagt ctcatctctg gaaagaaatg ctgtgtccag aactcagcag ctccatctat 420
tttttccagt cgaaagaaac tgatcttttag gcagttttta cttgg 465

<210> 146
<211> 351
<212> DNA
<213> Homo sapien

<400> 146
ccagccgggg taatctgtat gtggcggact tgagctacga cgtgggcggc aagtgcctgt 60
ttgaccagat cagcggcgtg aagcttatgc caactcatcg ttgataaat ccgaggatca 120
gttcaagacg tcgcagcggg tgattttggg aacgtcgttt tcggtcagta aattgtgggt 180
agcgacggag tggttgatcg gcaagaatga tccgtatatt ggcgggagca gctataccga 240
gagcctgggg gctgggggga gtaaccagtg ggagaatcag ttatataatga acattgggta 300
ctacttctga cttaatatct ccagcgtttt aactggcctt atcgaggca a 351

<210> 147
<211> 654

<212> DNA
<213> Homo sapien

<400> 147
acttattttt aattactgaa tattttcttag acgttttggg acagatttta tgtaatcttt 60
ataagtatga tttctgaaga aaagcaaagc cattagtagt tttgccttaa acttgtagac 120
taaaccaagt attgtaaaat aaacagcgat aacagtgata gtttttaact ctatggcat 180
tgtatcactc tggaaaatgt ggagtagctg taataaatct actcctgtat tatgctttac 240
agtgcaggtc ttagtttttc ttttttctca tttcttttga aatggcatct cgaacaaagt 300
ccaccaatcc ctttacaaaa gaatgaactg ctctctctgtg tgtacttcat agaagggtgga 360
atcgacaga ggcaggtag tgacagttat tcctgaaata caggagcaga gtacagtctg 420
ttgtggtttc cgggattccg cgcttagctc agccaattaa gcatgagaca taggccattg 480
agccacttag tagttatcgc agtggataga ttggtatgta agaggggaaag aggtctgctg 540
taaagaacaa cacttgtttg tctgtgggga aagaaaagca gaatcttgag atgaaagttg 600
gcatacaaat aggatactat cgccagtagg ttatattaca aaacatttat cggg 654

<210> 148
<211> 539
<212> DNA
<213> Homo sapien

<400> 148
tgaatatcat gagggtgatt ttcacctgat tgcaaaactg ccatagtttg aaacactttt 60
tcaattttacc agacacactc tgtcaagact tcatatactt ccaacttgca agcctgtgtt 120
ttgccttctc caacctaaaa aggaaaagct ttaaacgatg aacttacatt ctattaaacc 180
atcagacttg agcttatcca tctgtttagc gtgaatgtac aaaccaggta catttccacc 240
aaacacatag aaaaatcttg tgcatacacag ttcagctaag ggtagtagga caatccttac 300
aatcctcctt ggatttcttt tttaagatgt caaagaagca ggtaagcaac attgttcatt 360
tgttactggg tgttctagat caaaccttca caagctatat atatagcttc atatgctata 420
gcttacaaat ggggtaacaa agtaaaagaa aagaacaaat tatactttga cactttatag 480
tcaaagtata attaaaaaag aaatcctaca gtgggtaatg gagaaataga taatttttc 539

<210> 149
<211> 273
<212> DNA
<213> Homo sapien

<400> 149
tttttggtca ttctcctcaa ggagccgctg gatagtagtc ttgattgact tccaccttgc 60
ccctcatata gtccggtact aaggccaccg acatcccag gaacctccg aaccacgacc 120
gccaagcaac tcgaccacg ataggtgggg cctacgctct cgaagttgat tggatgctcc 180
cgctacagg ggggggtaca gaaggacgt catttgtgac tggacgcga agagctatac 240
tcagcagctt tcctctgtcc cagcccctag aac 273

<210> 150
<211> 200
<212> DNA
<213> Homo sapien

<400> 150
gtttttacta ccgtatggcc catttaaaag ggatgtgtac gccttacact ataaccctta 60
aaccacctag aaatatgaaa ctcaaactgc cactgacctc cctcaccaag ctccataaaa 120
gtaaaaaatt ataacaaacc ttattaacca aactgaacga acatatgggc gattgattca 180
ttgccccac aatcctaggg 200

<400> 151

<210> 152

$\langle 211 \rangle$ 243

<212> DNA

<213> Homo sapien

<400> 152

atttcaacaa	catactgtgc	gaggtagtta	taaatcttct	tagggggagg	tggtggtttc	60
tgttggaatg	ccaattttac	agcttctgct	gctgattcag	gttctttaat	tatgcttttc	120
tttgagtctg	cttcagatag	cacaacaaaa	aatgatgac	acttttcaca	cttgacaaaa	180
cgggtggatg	atacaaaagg	tctctacatg	tgtgcacaag	tcgccacatt	taggacagcg	240
caq						243

<210> 153

<211> 620

<212> DNA

<213> Homo sapien

<400> 153

tgtgtcttctc	taccttacca	tagccagttg	ctttcatttt	aaaccagagc	aagtaacata	60
ttagtgtactt	gaatcttcat	aagttaaagt	aaaaaacagc	aaaaaaccta	gatctttgtc	120
ttttagaaca	cagaccattt	tcaggaaagc	agttagctaa	gtgtttaatt	catgaatatt	180
gtatactgca	ccccctacca	caattttacac	aatcctgtgg	atagtcctac	ctcaccctgg	240
tcaacctaca	tgatccttaa	gctaattggcg	gatcacgatg	accttgtaga	catgcacaca	300
actatacctt	tgtccaacag	atcataatat	atctgctatc	caactggttt	tacctgcta	360
atcctactga	tttgggcact	gcttgatatag	tctctcaagt	tcacaggaaa	tgttgatttt	420
ctaaggtcct	cattttttaca	gagtatacag	gcaaagtgac	aggggaaaag	gaattagtct	480
aagagtaaag	ggatgattat	tatatattgag	ctaaaaccac	aaagtggctc	aggctttaaa	540
aaaaaacact	gtggataatg	acaaaaagca	taagtaaaaa	tattttgaga	aaaataaagt	600
acaagttttg	aacaccccc					620

<210> 154

<211> 843

<212> DNA

<213> Homo sapien

<400> 154

cattgttagt gaccaagta aatttatagt ttttaagtcc agaggaaaaa taaagcctat 60

tttttgtaa	cagtcttaat	aaataataaa	atggaataaa	gaaacccaaa	aaaaaagaaa	120
aagtttgat	gaaaattcat	ccctatctt	ttatcttgga	ctaagtagtc	aaatttctac	180
tatattaata	ttatgtaagc	gacacccatt	taaattcact	ctctttgata	gaaaggtag	240
ttgattatca	cacctgctat	tttttcaactg	ccaaaragac	tgcaataacc	tccctccatc	300
accctcaaaa	aacaaacaga	aaccatctga	ggcatagcca	ttgtttacat	attgtgtttg	360
tgtgcaccta	tctacaacgt	tctttcttct	aaggagttaa	tctgccaata	ttttcggtt	420
cagcagcagc	gctcttcttg	acagactaag	agaaggatct	acagaaaagt	catctgatta	480
aggttttggg	tcaaattaaa	actctctgga	cagaatctct	tttctctcac	ttggatttct	540
gcaaacagaa	agcagattat	tctctgga	caatagcgac	tctagaaacg	cttatgtttt	600
tcagactttg	gcagaacttg	ttaagaacag	catcatcata	atacatttgt	acaaactcga	660
atttcagtgg	ctcttttctc	ccacatgatg	catgatgaaa	tttataaagg	tctgttttac	720
ccccacagg	tcatttcttt	tgtgttccta	cagagccaat	aggcttcatt	taagtccaag	780
ttattatatt	aaccatccct	ttcactagac	tagagaactt	ctttttcatg	gtccatctcg	840
tga						843

<210> 155

<211> 674

<212> DNA

<213> Homo sapien

<400> 155

tttctgtgca	gccccaggtt	tgctccagct	attcacaagc	agaatataac	acaagaaaaa	60
caattcatat	cccttaggga	aaaaagagga	tcaattcatc	actcaatatt	taatacagcc	120
aaaatgagct	gcaaaaacaa	gcacacacac	aaatactgtg	aacagaaaaa	tacaagaaaa	180
tgactaagct	gggagctctg	acggggtagt	gacattgctt	aaagcactta	tcagtcccca	240
gaaaaaccaa	acaaaaaaca	ttttttacga	tgccatggcc	tcattggccc	ctttaaaact	300
gttgatggta	acaaagggca	gggggtgggg	agagaaaaca	caatcactgc	tccctttttg	360
ctgcagctg	tgactgcacc	cctcacggca	cggcatgta	cacaactacc	acacaaggag	420
gaccaagtcc	ctctgctggt	ggcctcctaa	aaggcaaggc	ttgagttttg	gctgatgagc	480
aagttctctc	cgttaccaat	ccctgccaac	cagcactacc	atggctgaat	tgatctaccg	540
ttttcctgag	taaactgtaa	ctggctacag	tttcggtaac	atggaaaaga	actcagctac	600
tacagccaac	tgcaataact	caggaacccc	ctccatccct	ggggctcctc	actcctagt	660
catcttgatt	ggat					674

<210> 156

<211> 671

<212> DNA

<213> Homo sapien

<400> 156

ccttttagtga	acacctttat	ctccatgtcc	ctcttagagc	ccagagagct	gcccataaggc	60
attttccaga	attcctcatg	tcacctagtt	caatttccat	taactcagat	cagccattgt	120
gattcaccat	ttgtcaggct	ctcaggttta	acaaaaccta	ctatcaccat	catccttcaa	180
cagccacagt	ctgaattgag	ccaacatctt	tttttctttg	agaaagaagt	gggctggggc	240
acaactttta	gtctgagggg	agctagtagt	cggcttgaca	attaaagcca	tccataacaa	300
cttttcttca	aatgtgttga	ctcctcaggg	gctaaactgc	tcttagctta	gaattatgct	360
ttactagaga	tctaccatat	aagtgggtta	atcactacca	tctgttaact	agttatatag	420
cttcagaca	tgaggagagc	atcaaacagg	gatggaagca	acccaagga	tatgcaagaa	480
gggcatgatg	aaccccttct	cctctggcag	gagaacaagg	ccaaccaagg	gacagactgg	540
aaagcactta	gatgtttaag	gaggagaaag	gggaagcttt	gaccagtcct	tgctttttgc	600
caagttcagc	cagttctccg	ctgcttgcaa	cctctagcgc	agtaacattt	tgagaattg	660
cagattttcc	c					671

<210> 157

<211> 474
 <212> DNA
 <213> Homo sapien

<400> 157
 cgcgttcttt aattctttta gcttagaaag tcctttacac tacttaccta aaggccccaa 60
 agtaaaacac acactagtag taaggctagt gcatttccct tctagcactc aaagaaagct 120
 taacattttt gacagtttgc aaataccgcc ttgtatttct gattcagcct tattcaaagt 180
 atcataataa aatattttatt aaatstatgt tgatctgcgt gcatttatga tctccagatt 240
 aacgttaggc ttctctgttg gcccctaact tggagggtgct tttttggatc cctcctcccg 300
 tgattcattg taatttcatt tcccttgta tggtctgac cagagaagat tctaaatatc 360
 tgccccaaa gccaaaatta tatcttttga aaagtgaat gaagagttga gtcastaatt 420
 tatttttagat attactgcct aaaacaattc cccaaaattt atggaagttg gagg 474

<210> 158
 <211> 584
 <212> DNA
 <213> Homo sapien

<400> 158
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 agtgccaagt ttagtcaatt taccctacct ggaatactat atacaactct gggctcctatg 120
 tgtgttaaaa tacatacagt gaagctgagg aagagccact gaagtaaaaa gtattgttta 180
 caagttggaa aggatgtaaa aataatctaa agtatactaa gtcaggaata aaaggcagag 240
 ttaataaaaa tgtggctggg actgatagac gaaacagata tattttctaa atcctggaat 300
 aattattaaa aaattttaca tgtatcaatg gattccagac tccatatttt aagtttcaca 360
 actactgtca tttaaaacta taccttattg aacgtctccc actctcaata aattacccca 420
 aatcactctt ctccaaaacg taaatttgga acacactgac ttacaaaatt tgggcttaat 480
 ttataggatg ttgtggccct caaaaatatc attgtgggct aaacaaaata aattcttgaa 540
 acaattctaa aaatcaatca ttgtccaaaa tgaacttttt ctaa 584

<210> 159
 <211> 671
 <212> DNA
 <213> Homo sapien

<400> 159
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 agatgaacca atccattgga agattactaa aattgtatct tccaatgcc tcctacagta 120
 agatttcttt ataattataa cccttggaaga caatttgaac tttattttaa tgttctgctc 180
 aaatctaaat ttcttctctc taggctgaag cctgatctaa ataaggaagt agttgggata 240
 tatccacagg ctgtcgaaca tggagctgca tctgagagac aggtggcagc aacccaaaagc 300
 aaagcagggg ctgagaacag gcaggttcca agagcaaaat ggaacttgaa agccaagtat 360
 ggttcactgt aaaggagaaa atatagaaat acggaactag aacacctggg ctgggatgtg 420
 gtaagcacco aaaatatagg aaaactgtat gaattcttgt gaagcagtaa actatgatag 480
 taatcatgtg acacatatga taacaaactc aaaacagggg aaagaggggc tttattcaat 540
 gctggagata agtgaaaaaa aaagtgaagt gtctcaagga cagaagttat catctcaaaa 600
 aggcataatca gctagatctc gcggaaacca tatgattatc ataattctag actctgttctg 660
 gtattacaaa g 671

<210> 160
 <211> 315
 <212> DNA
 <213> Homo sapien

<400> 160

<211> 607

<213> Homo sapien

<400> 161

<211> 443

<213> Homo sapien

<400> 162

<211> 686

<213> Homo sapien

<400> 163

ggtagacatc acctggatcc cccactctat tgcttacctt tttgttttgt aatttgatca 420
 gttcaagtta aaacaattta accaaaaact atgaatgttt atgatataat gaaatgattg 480
 ttaactttct tattgctttt tcacacacct ataaaagtaa ttttattact cccaagagaa 540
 atcactaaaag gcagaattac tagaggtaaa aataactagg gttggtacag tattactcag 600
 gagaagtcaa ggggagaaaa cttgtcccaa tgattcaaaa taattttggc atgggggggg 660
 ggagggaaaa aaatttggct tccttt 686

<210> 164
 <211> 706
 <212> DNA
 <213> Homo sapien

<400> 164
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 tccataacac agaaaatgca tggacatgca tctacagtag agttaaaaaat ttcctgtgac 180
 taaaaaatta aaaactggaa tcaccagtag caaatgtata gtcaatggct atgacaagaa 240
 cagatcctgc cgagctcata aatgcaatta ttggcttttt tgctttataa aaaagacatt 300
 acatatttta ttgcattatt ctccataataa aaaacatact accacgtagc tctccccatc 360
 cccattcttt gcttcagat ttttatagaa aataactgtt ttagtctggc cttggaaagt 420
 gaaccacca gcaccacct cacctactca ctcttcaatt caatatgcac atagcaaaag 480
 ccaacacttc aaatctcttg cccacatcaa aaaaagtagt ttcaggagaa aaacattaat 540
 accagttgaa taaaaataag ggcataaaag ctatgagaga gatagctctg ccatctgtct 600
 ctgggctaaa aatcaaggct aactattgcc tttggcacca caagggtcaa ggtccatggt 660
 tttattagaa aagtccccac aaaaaaatta aacccccctc acccca 706

<210> 165
 <211> 427
 <212> DNA
 <213> Homo sapien

<400> 165
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 aaattgtccc tgtttgcaga cgacatgatt gtatatctag aaaaccccat tgtctcagcc 120
 caaaatctcc ttaagctgat aagcaacttc agcaamgtct caggatacaa aatcaatgta 180
 caaaaatcac aagcattctt atacaccaat aacagacaaa cagagagcca aatcatgag 240
 tgaactccca ttcacaactg cttcaaagag aataaaatag ctaggaatcc aacttacaag 300
 ggatgtgaag gacctcttca aggagaacta caaaccactg ctcaaggaaa taaaagagga 360
 tacaacaaaa tggaagaaca ttccatgctc atgggtagga agaatacaata tgggtgaaaat 420
 ggaaaaa 427

<210> 166
 <211> 124
 <212> DNA
 <213> Homo sapien

<400> 166
 accatgtttt ogttgtgtgt gagcagggaa gggaactttc ctgccttatt taaacctggg 60
 ccgaggattc gtggaatctg cttgatcaga gactctgagg ccaaaaacgc atcatacttc 120
 ttgg 124

<210> 167
 <211> 232
 <212> DNA

<213> Homo sapien

<400> 167

tctgcatagc	aaatatgatt	taagaattta	acatcattat	ttgatcacia	gcgtaaatat	60
gtcaccataa	ataaatgtaa	attcattgta	caaaaattcc	caacaactct	taatacaaat	120
atggtacatt	tgacagtttc	tgaaacagat	tattttttaa	acttttttaa	acctaagctt	180
tatttttttc	ctggttatta	gacacacaca	aaaaaataa	aaagaggctg	gg	232

<210> 168

<211> 677

<212> DNA

<213> Homo sapien

<400> 168

tttcacaatt	aaccaacatg	caaaaattct	cagactaaac	actgagaaat	tcttcataca	60
atgcatttgc	caccttattg	cattttttaa	atctttattc	tatagtgaat	tggtattccc	120
aatctgccta	agcaaaggca	tgcccttcta	acaagatttg	cttagagcag	aggtgataga	180
aggaagaatc	cgaagaccct	ctggcatggc	aatctgggag	cagcacattg	ttgatggagt	240
ccaagtgagc	acatttcaca	caattcattt	agtgacaagt	gggcttgctc	ccttttcatc	300
caggaaaaaa	actactcaca	gaccactgcc	cagaatctgg	aataagaacc	ctcattttta	360
ggtattcttc	ccaacaaata	aatatctaaa	tattgaaaag	gggcatatca	gaaaacttaa	420
aagacacaat	aaccaaaaacc	aaaaccctct	tcaaaacaag	taagcaatgt	ctgtatttag	480
ttcactctaa	aacattctta	gcttttcttg	cagtttggtc	ctaaaagatt	tgattgggca	540
caagaggaac	gaaattatta	ataaaaataa	agcttatttt	tgtttttgct	gtggataatc	600
ggtacaaaac	gtttccagat	ctgagactta	aatggatctt	ttaaggtgaa	aaggagaatg	660
ccaggttcta	ctgaaat					677

<210> 169

<211> 635

<212> DNA

<213> Homo sapien

<400> 169

ttaagaagac	tgggcattta	tactctctct	tgctagtcag	cctggagcaa	gcttggagca	60
gacgcacatt	tttgtactgg	cacatattct	tagacgacca	attatagttt	atggagtaaa	120
atattacaag	agtttccggg	gagaaacttt	aggatatact	cggtttcaag	gtgtttatct	180
gcctttgttg	tgggaaacaga	gtttttgttg	gaaaagtcgg	attgctctgg	gttatacgag	240
gggccacttc	tctgcttttg	ttgccatgga	aaatgatggc	tatggcaacc	gaggtgctgg	300
tgctaatact	aataccgatg	atgatgtcac	catcacattt	ttgcctctgg	ttgacagtga	360
aaggaagcta	ctccatgtgc	acttcctttc	tgctcaggag	ctaggtaatg	aggaacagca	420
agaaaaactg	ctcagggagt	ggctggactg	ctgtgtgacg	gaggggggag	ttctggttgc	480
catgcagaaa	gagttctcgg	cgggcgaaat	cacccctggg	tcactcacat	ggtacaaaaa	540
tggttttgac	ccgctaccga	cagatccggc	cgggtacatc	cctgtctgat	ggagaggaag	600
atgaggatga	tgaagatgaa	tgaaaaaaa	aaaaa			635

<210> 170

<211> 533

<212> DNA

<213> Homo sapien

<400> 170

ctgtgatctc	acaagtgtga	aaaatcttat	gaatgtaaaa	tgtgtggaga	ttcttctttg	60
tttttagctt	ccactttggg	aacatgtcaa	agcacacatt	gagaagtccc	atgagtgaag	120
gagatgttgg	aaagcccttg	aacttggtcg	ttaggaaaca	tccacactga	agaggaacct	180

gactgtatgg	aaggtcaaaa	aggctgtatt	aatttacatg	caaaaagtca	cactagagga	240
atgccatatc	agaatgcttt	tggtaaatat	acatgtttta	aagaggttat	atatcattaa	300
taaaaatatc	tagctgggtc	gaagaccctg	agttatctca	attgttcacg	gttacagatg	360
gaactcttta	ttattgagga	gttccactct	ttccccatt	tgctactact	acacttccct	420
agtctttaa	acaatttttag	gctgggtgca	gtggctcatt	cctgtaatcc	cagcactttg	480
aaaggccgaa	gcgagtggat	catttgaggt	caggagtctg	agaccagcct	gga	533

<210> 171

<211> 568

<212> DNA

<213> Homo sapien

<400> 171

cccttgsc	aa	actttccctt	aagtattgca	ctacaagtct	aagacacttt	tcactcaaag	60
ttccttcc	ct	ccttacctct	cttttaactt	ggagtcagac	tttcatcagt	ctgacaactt	120
ctccctgt	ct	ccttcccttt	cccccttca	caagcatttc	acctaacaaa	tttcttatgt	180
gcttaatcc	ct	cttagaag	cagatgccaa	gatgggatta	agcacataag	aggctcctgga	240
ctaataca	at	gacaaaggct	ccccttgaag	catcacacta	aaaggaaaaa	aaaaaaaaaa	300
acctagcca	tt	tacattaa	ctatttctaa	aatatagtat	ttgcttccct	atttgctaaa	360
acaaaatata	ct	aaacatga	ctattccaaa	aatctgtagg	gtactaagaa	tatgaagaga	420
ttcactctac	tt	caggggat	ggagttgtag	tagaaaaggc	tttgtggagg	gaggggtggtg	480
tttgaaatgt	act	tttaaaag	ccatcctcaa	agcctcgagg	gctataacctg	gcctggtgat	540
tatccaagga	cag	tccattc	aaacaggg				568

<210> 172

<211> 167

<212> DNA

<213> Homo sapien

<400> 172

ccattttac	ag	gaatcagcca	cttcagttca	gacagcttta	ttaaaccgcc	tggagcgaat	60
tttogaag	ca	tgttttcctt	ccatacttgt	ccctgatgct	gaagaggaag	ttacttccct	120
gaggcact	tg	ctggaaacaa	gcactttgcc	aataaaaacg	agagagg		167

<210> 173

<211> 391

<212> DNA

<213> Homo sapien

<400> 173

cctcccaa	ag	tgctgggatt	acaggcatga	mccmccmcgc	cctgatgata	gacacgtttt	60
taactttc	taa	aaatatatga	tcattgattgt	gtctgtggag	acttgacat	atactaaatt	120
ttaamcaa	att	agagatattt	gttcattacc	acattttggg	agtcattatt	tcctctatga	180
agagagaa	ag	gaatttgata	caagttcaca	ggggcttcca	gtagattgag	acttttat	240
ctagctga	gc	tgctgatgta	tgaatttttt	ttgktattat	gactttcata	tgtattaaaa	300
ataaaatg	aa	aaaacaagg	attaggtgag	gaacctatac	gtctctaata	tgcaaaatac	360
cacagaaat	a	tgactgktg	ggaaaattag	g			391

<210> 174

<211> 474

<212> DNA

<213> Homo sapien

<400> 174

gaactcagag	agaggattgt	cacccttggc	atctgagctg	acactataag	gacaatgagg	60
agtctccttg	gggatatag	gggagatgga	aggacgatgc	ctgtcctacg	gggtcttggg	120
aggttaggga	tacacactgt	gagctgccac	aggctcaaca	gtacggatag	ggggtgctgg	180
aaccagccag	ggctctgatc	accaagctat	gtgccccatg	cagaggaagg	ggtagtggca	240
caactgaacca	cccagccaca	aggctatctc	cccatacagg	gcacctttaa	aaaaattatc	300
cttacagggg	aagacgggga	ggaaggatga	actgtgtgcg	gtgatgttgc	agtgagtgtg	360
agtttgtgtc	cgtccgcttg	tatgagggcc	taccttttac	taactagccc	ccaactttca	420
ttatctcccc	tttttctgtc	tacccttctg	ccttttttaa	gtggcttgca	atcc	474

<210> 175

<211> 655

<212> DNA

<213> Homo sapien

<400> 175

ccttgcaggg	gtggggatgt	gtgggcttgt	tcaactgttac	agcccatgta	tacctgaagg	60
gcaacatgta	cccacaaatg	ttccaggagg	taaataaaaa	atacaattca	gcctcttcta	120
aaccatcctt	gttgatatct	ctgctacttc	cgaaagttaa	ttcgttatct	ggactccata	180
atttttccta	ttaattcacc	ctatgtccaa	ctccaacagt	gaaaaaaaatt	tatttaattc	240
ttgcaataag	cctataggca	ggcagcatta	tcctcagtct	gcagataaag	taaggctcag	300
agaagcttgt	atactgtcac	ttaggtagta	attgcaagag	ctggcattca	gaccagact	360
gtgggactcc	tcactccatt	ctctttcccc	ccactaggct	gctccttaaa	atacaatgga	420
tgtttgatga	acgcttggg	gaatcctggg	tggacacagt	tccttttcgg	ccaaaagcac	480
cttgacgact	tgtgaagaat	taatctggaa	aacttaacct	atttataaaa	acgtgttatt	540
aagggcaggt	tattcccacc	ccctttacca	aagaaacccg	ccctgacctt	tttttactgg	600
gggttggtct	tgggcatttt	caacaagggg	ggaacagttt	aaaaattccc	ccctt	655

<210> 176

<211> 660

<212> DNA

<213> Homo sapien

<400> 176

cctgggtcaaa	gtgggcatta	ccattcaagc	attactagac	atcacccgtaa	cgaaggctct	60
gttcacatga	aactacccct	tctccattgg	gggctcagac	tctgctctca	tccaggatcc	120
tgaactctgc	tccaggcacc	tgttcaacct	tctctcccac	ccactgcctg	tcacttcaact	180
gactccagtt	acattgaaac	aattttcagt	ctaaggagg	attttctacc	tttcagagct	240
gacctccgac	tttaagactt	gacaggat	tatcttgaaa	ccagagagg	agctggagg	300
aaaaaaaaact	gagcaagcac	atcaatgcct	tttccaccct	tcttcaccc	ttccacactc	360
accgactgcc	attaccaaaa	cgccaagcac	aaccggtttg	gaacaagacg	cattccgttt	420
taattaaaac	caactcatta	tgtatttttag	tgggggggaa	gggggggcaca	atcagggttt	480
tcaccaccaa	attttccaca	cggttttctga	acaccattgc	cttttaaaaa	actatttttc	540
cacctccaaa	atattttatt	aaattttatt	tattacggag	gtgggtattct	tcctttggga	600
gccaaattgg	gaaatttagg	gaaccttttt	tattaccgag	ttttttgggc	gggtaaaccc	660

<210> 177

<211> 459

<212> DNA

<213> Homo sapien

<400> 177

ctttttctct	tcctctgtgg	aatgggtgaaa	gagagatgcc	gtgktttgaa	gagtaagatg	60
atgaaatgaw	tttttaattc	aagaamcatt	cagaamcata	ggaattaaaa	cttagagaaa	120
tgatctaatt	tcctgttca	cacaaacttt	actctttaat	ctgatgattg	gatattttat	180

tttagtgaaa	catcatcttg	ttagctaact	ttaaaaaatg	gatgtagaat	gattaaaggt	240
tggatatgatt	tttttttaat	gtatcagytt	gaacctagaa	tattgaatta	aaatgctgkc	300
tcagtatttt	aaaagcaaaa	aagggaatgg	aggaaaattg	catcttagac	catttttata	360
tgcagtgtac	aatttgctgg	gctagaaatg	agataaagat	tatttatttt	tgktcatgyc	420
ttgkactttt	ctattaaaat	catttttacga	aaaaaaaaa			459

<210> 178

<211> 720

<212> DNA

<213> Homo sapien

<400> 178

ctgcaagctc	ccactccttc	catttatctt	aacgcccagg	ctgacttcta	agctgctttt	60
cactttccta	cctccactgc	attttcgccc	ctgataatth	ttgtaagctt	acctaagcct	120
cccttctttt	gagatcccct	tcttaaaagg	gtccattcta	ttaacctac	cccatatcca	180
gttactttta	ctacctgctg	atctatcgct	accttggtcca	attcatggga	attacagggg	240
gcactgggac	aagagtaaaa	tgatccaaca	aacataatgt	tgcatthtaa	aaaataagct	300
aaaagatact	gatgactttt	tataactaca	acataatcgt	ttgtgaataa	gaacatatat	360
agtaaaaaga	tgaaaatgtg	aacaggttga	ctatttccta	aatttatggc	agaaggttgt	420
tctggagagg	atgggaagaa	aaaatgaagg	ctggcagtga	tgggtgggga	aatgcaacct	480
ccaaaattat	ctatctatat	atttttatta	aaaacaccca	cagtaattat	ggcaaagtgt	540
aatggtttgt	ttgttctaag	gttttgata	catttaagat	ctcttgcttt	ctgggtacca	600
tttcttttct	tttcttttct	ttttttttca	aattaattcc	aaaagactta	tatctgctac	660
atgaagaacg	aagcaagttc	agctctcttg	gctgaaatgt	tcaaattgctt	gagggcaagg	720

<210> 179

<211> 427

<212> DNA

<213> Homo sapien

<400> 179

ctgtgaatct	gtctggttct	gaacttattt	tttagttatt	ggcaatcttt	gtattactat	60
ttcaatctct	tcctggttta	atctaggagg	gttgtatatt	tccaggaatt	tatccatctc	120
ttgtaagttt	tctagtttat	gcacataaac	gtgttcatag	tagccttgaa	taatcttttg	180
tattttctgtg	atatcagttg	taatatctcc	catttcattt	ctaattgagc	ttatttgaaa	240
cttctctctt	cttggttaat	cttgctaattg	gtctatcagt	tttatttatc	ttttcaaaga	300
accagctttt	tgtttcattt	atcttttgta	ttgtttttgt	ttgtctcaat	ttcatttagt	360
tctgctctga	tcttcgttat	ttcttttctt	ctcctggggt	tgggtttaga	ttgttcttgg	420
tttctct						427

<210> 180

<211> 728

<212> DNA

<213> Homo sapien

<400> 180

caaacacaaa	agtcactgtg	tgtgtgatgc	ttctccaatt	ccactcatcc	tggctgccat	60
tcatgcacta	gtgcatgtat	gcattttttac	attttttaaa	ttacaaaaat	caacctatta	120
taactgctta	gatatatatg	aagtaaaaaat	gaaagttctc	cctttacatg	acccatcccc	180
catcatttcc	ctcttttatct	tatactgtca	gcattcccag	cttgtagcac	agtgtctggc	240
aatagtaaat	octcaaaaaa	tgatcaatga	ataatttaat	aatgattaat	aaataaatta	300
atgatgatgg	tgaagataaa	ttttagcatt	tattgaacgc	taactacaaa	ccagggagtg	360
tggtaaatat	tttataaaaa	tcaatgaatg	agctaaaaatg	ccattctatt	atttttttgg	420
atacggttta	atatttttact	cataaatatg	cttaaagaat	attataatta	tatgacttag	480

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aatggtaaaa caatatgtac agcagtatcc ttttttttag aataaaaaata taaatatgtg      540
ctcacatatg tggttggggc atgcctagaa acccgattag aacgggattt tttcttacca      600
ccattttttt tacctgggaa aaatatggga aaattttatt tcccttcttt ttgggttctaa      660
aatttatata caggagccta tttggctttg gataaatcat tttaaaaaag gtgggtttaa      720
aaaaaaaaa                                     728

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<210> 181
<211> 546
<212> DNA
<213> Homo sapien

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<400> 181
acaatccttt ggaagacact actgggcttt ggggtgctgct ttttaataat tgagttatatt      60
tgagcttgcc aagtaggatc tattgcctgg actaaaattt atttcctaatt cttctgatga      120
ccaagaaagg aaaaattaag tttgcagatg ggagatgaaa tatagccagc gaatatgcat      180
actggttctg aatgaaagga attaactttt cagtcaagaa acagtctgca tgccgtaaat      240
tgaatttttc ctgcaactgg aatgattggt taattctttt tgaacactgg cctttctccc      300
caagaacact aatgaattgc taatattttt taaagaaaaac tgggtttttt attaggttaag      360
ctccacttcc tcttattttt taatccctaa agaaaactgt taaaagggaa tggatctatc      420
acgccttttc ttttaaaacc acctttttaa aaaaggattt ttccaacccc caatttgctc      480
ttatttttaa attttgaacg ccaaaagaag ggaaataaaa atttttcctt taattttacc      540
ccctta                                     546

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<210> 182
<211> 333
<212> DNA
<213> Homo sapien

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<400> 182
ggccactctg actgggtctg ctaattcaca tgctctttgt gacatacggc tctaagaggc      60
agaggctgga agagaagtat gtgggtttgt ggatcaagat acccaagttt cagtcttgac      120
actgctatta cttagtcagg tgaccactgt aacttcatct tgattgagcc tcagatgtct      180
cacctgcaaa atggagtttg aaatttgcta tggttgggtg tcacacggat taaatgaaat      240
aatgcctgtt aagcgcctat ccagcactta ataagatggc cactgcatca taatgctttg      300
ggcacaagta acacaacatc caacccaaag ggg                                     333

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<210> 183
<211> 393
<212> DNA
<213> Homo sapien

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<400> 183
ctgaattttc tgggctttat gtggcagtgt ggtaaaaaata tatgatcaga tttcactgtt      60
aagaaaaattc tttcagcaat acatgtagag tcaagtttct tgcattggata actgaacatg      120
tgggttatga gatttttaaaa aatgtctcgt gacaaacttt acggaaatgc aacaatctgg      180
acatctagtt ttgtctgaga gtggcgtgga tatgaagaac tgtgctgttg gtgctgatgc      240
cacactaagt tttggcagtc acactcttgg ttcttcatat ttgaggagat gggatggtga      300
ggaggcctgt tggctttatt ttattacgtg ccaccatcta gaatacagat tcttgatat      360
ttcatcttca caaaggtgaa gctgcaaact cag                                     393

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<210> 184
<211> 700
<212> DNA
<213> Homo sapien

```

<220>
 <221> misc_feature
 <222> (1)...(700)
 <223> n = A,T,C or G

<400> 184
 ccaggscawt gaggaaaaagr gaaagaatwt arrggstwt caaataggaa aaraggaagt 60
 ccaaattggt ccntgttkg ccagataacc atgattgk gk atttagaaam ccccatgwt 120
 tcagcccaaa atctccttaa gctgattaag camcttcagt aaaktctcag gataaaaaat 180
 caatgtgcaa aawtcacaag crttcctatm cgamcaatam cagmcaaaca gagccaawtc 240
 atgagtgrac tcttattcac aattgctagt aagagaagaa aatmcctagg aatacaactt 300
 mcaagggatg tgaaggwtct cttcaaagaa gaactacaar ccrctgctca aggaaataag 360
 agaggmcmca agtaaatggg aaaagcattc tatgctcatg gataggaaga atcaatcccg 420
 tgaaaatggk gatactgcc aaaataattt atagattcaa tgctatcccc atcaagctac 480
 cattgacttt cttcmcgaa ttnggaaaaa tctactttac acttyatagg gracaaaaa 540
 agaagccwt gtagccaaga caatcctagg caaaaaagac caamcctgga ggcatcacag 600
 tmcytgactt cmaactatwc taccaaggny tmcrgkgmcc aaaacagcac ggkacntggt 660
 mccaaccrg acwtwtwgac cmmcagacac agaacmgagg 700

<210> 185
 <211> 192
 <212> DNA
 <213> Homo sapien

<400> 185
 ccagyctttt ttttaagtaa ggcgtttttt aagctcattg tagctacaaa gtcaataaat 60
 tgggtctttgt tatttttacc tgaaaaggct gttaaagggt aaaatgacaa actcaaattc 120
 aaagggattg gaggatttgg tgtttatgat ttctcagaac aacaatctag agaccaccag 180
 ggtgggtttc ag 192

<210> 186
 <211> 688
 <212> DNA
 <213> Homo sapien

<400> 186
 gtgctggaat tcgcccttag cgtggctcgc gccagagtg gatatttctt ctggatagat 60
 ttcagatagg tagttccctc aaataagatt atatgggtt gcattttcaa ggcagagttg 120
 tatacttct gctctttatt taaataaaaa aacttgaaaa tctgttctgc ccagtattgt 180
 aagcgctcag gtacaaatat gaatgaaaca atctctgcct aagtaacaca agtataggga 240
 caagattctc agtaaaattc tcacgtgaaa tttgtaactc actagacact atcaggagat 300
 caataattat gtaattaaaa aaaataatta cctgccaaac tgggttcttc tttggcactt 360
 ctgcttggtt ttaagacaat tctcacatag aagcttatta ttccccatta gtcattccat 420
 agatgtaaaa ctggtagaaa caggacttga attgaacatt ctttacaagt aagttatata 480
 gcttctgaaa aaagggcttg aaaaagcatt tttggggact ataagaacct tcaaatgctt 540
 tcccccttta acaaacctta aaattatatt gaaaataatt taagggggct gattttctct 600
 tgtaaaaatc ttgaacccca cttaccaggt ggttggtcaa accaaagttc aaaaaaaagc 660
 ttctggcctt tcttttatcc cacttgca 688

<210> 187
 <211> 779
 <212> DNA
 <213> Homo sapien

<400> 187

<211> 394

<213> Homo sapien

```
<221> misc_feature
```

<223> n = A, T, C or G

<400> 188

<211> 681

<213> Homo sapien

<400> 189

ctgac ttggtctat aaaacagggt tattggctgt ggctgcactc aatatctaaa 60

attag gaagtgcctc gttattgtca ttaaagatat ctaaatatgg tagaccaaag 120

tgaga aacacatatt atggactgag ttctgtttct tctgctgtgg cgcacctaag 180

gcctt ccttctctcc ctccccttct ggccggcatg gtatctgagc tcacagacag 240

gcatg ttagaatcat cagatcatga gcaccgtgct gggatttagc cctctccaaa 300

ttctt acagtcata ctttgcttaa atcctcagtt gttgaggtct gctctgctgt 360

atccc agctataaat ttcccccaa tgtggggcct agataaagta gaaggtggat 420

cagct tattttcatg ggatgacagg aactggaaag agaaagggca ttgaaaataa 480

tattc cagaatagca ttaaccctct tactgttcaa gaattaagaa agcctactta 540

gaggg ccttgagaat gatacccaaa tattggtctt tctaccaaaa aatggccttt 600

tatct gctttcctgt tccccaattg gctttttaag tagaattaag ttacctaataa 660

cctga aggggtggttt t 681

<210> 190
 <211> 839
 <212> DNA
 <213> Homo sapien

<400> 190
 caaatacatg atttccattg gcatagactc ttctatagtc tctcaggcac accttatgac 60
 taataagaac actgtcttct agatataagc caagttttag gagttatctt tgtagtttct 120
 gtgttgagac tatgggtctt ccctgtgcaa agacttgatt agcaaatact atttgaaacg 180
 atcccaaatt catagtgcag ttgaccaccc ttctgatcaa ggggatctct gtatatccca 240
 tgaaagcttc ataggtctca ccctagatta agtgcttcac ttctcaagac agtgaacaga 300
 tggaagactt ttgtagttat cattatacaa ctgtgccctg tgtgttttat tataacaacca 360
 gagaactgag gcactggctt tacctgtcag ctacgccagg ggtgtgacgt catctttctg 420
 acttgatcac acatgccaca ttgcttaata tttcaagctt agactgaaat aatcctgtgg 480
 taaaaaattt ttggggggct ggggaggtaa agaacaaggg ggggaacttt ggaatatattt 540
 tattcattaa tcatatttcc cgaattgtat tttattttga aatgaccata agggacttaa 600
 atacgtattg tggttaaatt aaatggaccc aaatggaggt aagtaaacct aatgggacaa 660
 atgaataaaa ggtttatgac tgggagcatt taccatgaa cctccttaga agctatttaa 720
 cctttctttt ggaaagccct gaaggctggg aacttaaatt ttaaagacag tacctatttc 780
 cagaatcgct tccaaatggc catgttttaa agggccaaca ttttgggatg gccctgccc 839

<210> 191
 <211> 697
 <212> DNA
 <213> Homo sapien

<400> 191
 ccatactgaa tactgatttt ctaatggaac tctattcaat ggcgattgta aaacctgag 60
 gctccgttac tattatggag catactttca tctcattctc ggctattggg caatatgtat 120
 ctcataagat tttatcacat ttcacagatg aactgttaat tgattccatg ggtacgatta 180
 ggcgagatcc aagctggagc tgcagctctg agtcccataa attctttgtg cttctgtaaa 240
 gaataaatct gtttttaatg caaattaaaa ctactggcag ggaatttttg cttccagtta 300
 ttaaaagact ggaaatgtgt aagtggagaa aggcaataac tgcagtaatc tcttaccgga 360
 ctctattata attccaaaca tacataatgg tgagaaaaac cgggaaggga agaattgtggc 420
 aatgtccact ctttgcccca aacataaccc ttaattttcca tggcgggccc aaacactggg 480
 aaaaaccaa atggtaccct ctatagcatg caacttttat ttcactccaa acgaaaaatt 540
 attttgacta tggttgagg aatccattag tagaagaagt tttataacct ataggaaccc 600
 ggccatttca tttctaccaa atcacaggaa ttttagaatg ggcaaggaat ttacaggaag 660
 acttgcccaa ttatcttttt ttgggggact aaaccaa 697

<210> 192
 <211> 687
 <212> DNA
 <213> Homo sapien

<400> 192
 ctggttacta tagctttgta gtataattta aagtcaggta atgtgattct tccagttttg 60
 ttattttctgc ttaggatagc tttggctatt ctggatcggt tgtggttcca tataaatttt 120
 aggatagttt tttgctattt ctgtgaagag tgtcattggg actttgatag ggattgcatt 180
 gaatctgaag attgcttttg gtagtatgaa cattttaaca atattgattc ttccgattaa 240
 tgaacatgga atgtttttcc tttatttggc gctctcttta atttccttca tcagtggttt 300
 ataggtttca ttatagagat ctttccttct tttgggtaat tcctacgtat ttaatttatg 360
 tatcgctatt gctaaatgga atgacttttt aaatttcttt ttcacattgc tcctgggtggc 420
 atattaaaaag ctactgatgg atggtgattt tggattctgc cactttactg gaattgggtg 480

```

atcagttcta atcgttttct tatgcacccc tttaacggttt ctacatgtaa gaatatatca 540
ccttcaaaca cggataatth gactttcttc ccataccaatt gggaggccct ttatatcttc 600
tcttggcctg aaggctctac ttaaaacttc ttatcccttt gttggaataa cagtggggac 660
aatggacat cccttgatcat ggtccca 687

```

```

<210> 193
<211> 493
<212> DNA
<213> Homo sapien

```

```

<400> 193
ctgctaaaat gatgttgcta aagcattcct ttttcttttg attaaacttc atgtttacaa 60
aaaaattaat tctagcagaa taacgaatgg ttttgttttc tagttctctg ctgaatgaac 120
agttttgcca attatcttca tagagtagtg atataatgaa tgcaacctca aatgcaaacc 180
aaccaattca cagtccatac cccaatcact tccttcatca gctcaaaaaa tcgctaagtg 240
aaccagtaga atggtttttg agcagtaata ggaaagcaaa tagaaagtca agggggactt 300
tcaacgccaa caagaccaat tcagatcctg atctgactgg tttctaatac aatctctttc 360
cagagtaatg gagcatgagt ctgccacaca gaactttaga gagagtcctt tatttcaaag 420
actgtaaagt tggaagaatt cattcatctg caaagtcaaa tgtcaaaaagt tgtgcttccc 480
actcctcatc agg 493

```

```

<210> 194
<211> 424
<212> DNA
<213> Homo sapien

```

```

<220>
<221> misc_feature
<222> (1) ... (424)
<223> n = A,T,C or G

```

```

<400> 194
cyagggcant ttagcangas aaggaaatan mggggattca attaggaac wraggakarw 60
caagttgtcc stgtmtgcag atgmsgtgat tgtatatcta gamcacccca ttgtctcagc 120
ccaaaatctc cytaagttga taagcawctt cagcarmgtc tcasgatscr acmtcwatns 180
gcraaantca cmwgcattct tatacaccaa tawcagacaa acagagagcc aaatcatgag 240
tgaactccca ttcacaattg ctacnmaaga gaataaaata cctaggaatc caacatacaa 300
gggatgtgaa ggacctcttc aaggagaact acmaaccact gctcaaggaa ataaaagagg 360
atmcaamcaa atggaagaac attccatgct catgggtagg aagaatcaat atccgkgaag 420
atgg 424

```

```

<210> 195
<211> 229
<212> DNA
<213> Homo sapien

```

```

<220>
<221> misc_feature
<222> (1) ... (229)
<223> n = A,T,C or G

```

```

<400> 195
tgaacaccct tnggaaggaa cctgctcgna tgtannanaa anggaccgga cagtctgcta 60
aatcgccct ctttagacgc ggcgcgcgcg ggcagagttt ttctctggtg ctttgacctg 120

```

tatttggttt aatgggtttg tcctaattctc ttcaatcaat aaaattgtgc gtattttaact 180
 aaaaaaaaaa aaaaaaaaaa aaaaaaaaaa aaaaaaaaaa aaaaaaaaaa 229

<210> 196
 <211> 557
 <212> DNA
 <213> Homo sapien

<400> 196
 gcggtggctc atgcctgtaa tcccaccact ttgggaggct gaggtgggca gatcacttca 60
 agttgagagt ttgagaccag cctgggcaac ataacaaagt gagatcttat ctctacaaaa 120
 aaattaaaca aacaaaaaaaa caaatcaaca ttcatttgca gggctctttg gtcttcttaa 180
 agaacaaaca tatgaaataa ataagctgat tcttaaagat aacaaatata atgagctttc 240
 tcaactgtaa aagcatctct aagttgttct atcaatgcat atccactcca tgaactaacc 300
 tgaagaaagt gttgaccatt ctacccaatt aactgtaaac taagattgct ttaatgggtt 360
 gcctaaattt gagtaccttt aaatttttgc tttttatcca aattcattct cccttcttca 420
 aattaaatag ttttgtaga aatcggataa gcaagatgta ctttttagaa agggcaatag 480
 aatcctacaa catgctagaa tttgaaatgt ttttttaaata cagtmmtttc tctatgctag 540
 taactaagaa aattata 557

<210> 197
 <211> 624
 <212> DNA
 <213> Homo sapien

<400> 197
 ttttactacc tatattttaa atgatccctg acgcccctca agacaaatat attaattttt 60
 ttactttgtg ggatagagat cagaaaaaga gtagagatga aaatactgga gaaacaatgc 120
 aggagatatt tatgaggtga gaatgtcaag aaacttgtaa agggagaata ctataatgac 180
 ccctgaagag agagcttttag accagttgag tattagaggt tgccacgtgg ctattcatcc 240
 actaataaat acaagaaatt actaaaatgg aagccactgg aaatatgttt tgaggaaggt 300
 gagaatgtgg acctattata aatgggtgaa tatgatttct ttctcattaa gttcataaat 360
 aactttcaga catgtaacag tttatgaagt gtgccgtagt catttagtat aagtttttata 420
 cacaaaagtg tttttactaa gactgtcaca ggttcttttg tgaatcttgt ttgtttttcc 480
 tcattgtaaa tactgcaata gaacatttgt gtcttaacat aaggcaataa atgaccttaa 540
 gaaccttcac ttttatatag aaagtggagg aaaagtgggc agagtaattt gttgattata 600
 gataaaagct cttgtagaaa ttgg 624

<210> 198
 <211> 175
 <212> DNA
 <213> Homo sapien

<400> 198
 tttttttttt tttttttttt ctaacactta tgcatttatt ttcattgtgta agaagaaaaa 60
 cgtaactagc acgtgaacat gactgcatgg atacacggct cagcacgagg ctaaagtcag 120
 aagtgagtga aagcaaaacc gcatgttgat ttaagtgaata taacagaaca gaaaa 175

<210> 199
 <211> 871
 <212> DNA
 <213> Homo sapien

<400> 199

```

ctgttgatca atgatgagct cccaagagta accagcctct atagatcag catcactggt      60
ttctcaggaa aagcatcacc attgttcato ttgctgcaaa atgtatgcac aagtatcttt      120
ttatttttaa aaaagccctg acattttatg actgctgctt ttctaagata ttttcaaata      180
tacagtccat acgggttcaga cacaatggac tggggataga gacggctata gtgccgataa      240
tggagaaaact agccagagct tcagatatat gttttccagg acatctcaat aattgggtac      300
acctcacaat atgtgagact tgacgtcgag tggcacggca tactctggcg caggcacttg      360
ataaagactg tgtttgcaaa tacttagcct gcacttcaag ataccaggca tctaagcacg      420
tcccagatgg tgacagttaa tcttcaaaaa accctatgtg gaagtattat cattgtcctc      480
attttacaga tgaggaaaaa gagacacagg gatgtcaata tcttcctcaa ggtcacacag      540
caagtaagtg atggaacagt ggctcagcca tgaagctatt gctgttaacc actaggttga      600
tttgccctta ttaatttctt cctaaaaactg cacatttccc gttagtccct ctttttggtc      660
tgtcgtttga ctcttggcta ctgcttagag gaagattcat tctattattt tctaacttag      720
taaatatgtg caactccttg gggacatgac caggcaaaaag ctggatacag aaatgtatgc      780
ccaaacacca tcccaagtta cccctaacag gtcttttctg gacctgttt gtaagggggg      840
tatatttgga aaaattttta aaattttctg g

```

```

<210> 200
<211> 737
<212> DNA
<213> Homo sapien

```

```

<400> 200
gacattttga aggtaacagc aatatctgtg tatagatggg gttgtggttt tgttatttat      60
ctgctattgc tgaactatcc tttgtcttga gcgataaaaag agaagtaaaa tactaaagaa      120
ctgaactgtc catttctgga ccatgagtaa agatgctggc tgtcaaactt cctgttcata      180
cattagttta tttatagagt gtactctcta tgtaaggat tgactgataa tgttactttg      240
acttcagata gcttgcagtt taatggagga agaagacaaa catgcaaata actagggtcaa      300
tgaggcatcc tttgtgttcc attggaagct aggctgcttt gtaaccttgt taatttctgt      360
gggtttggag tgcattcatt agcaaataca cccctgttcc ttatccattc tctgcttttt      420
tctttatttg gcatttgatg acattttttc atgtggggaa attgagtcag gtgaggtgga      480
aagaaaataa ggacacgaca ctaaaattctt tgatgttttt ccttaaaaaa ttgtttttca      540
agtgtcccat aaagggttgt gaagttttta gagccatagg acttggatta ttgtgaaaga      600
gtgtctctag ggggccagggt taaaccattt caaggactct ccttctctca tctcccttgt      660
tccaccaggg gtggcgaccc ccaaaaagca caaagcctcc ctttcttcat gggaagggta      720
aggaacggaa ggaacc

```

```

<210> 201
<211> 493
<212> DNA
<213> Homo sapien

```

```

<400> 201
tctagaaatg cagctttttat ttattacccc atttctttca agtccttgga aaataacata      60
ttaagggtac aagaaattaa cacatgatgg aaaagtcatt gtgacgccaa tgaatttcat      120
tgagtataaa ctcatctact tcaaatttat tttataacac aacctaagat actcaagata      180
attatttaat ggttagctct taagttgaat tgggtctacat aatgcgtggg aagaaaacca      240
gatttttagc cttcttgcca aatccagacc tctggttgat ttttctttga cagaagatgc      300
aagttatttt ccaatttcac aattaaatgt atttaacatg aacattattt tgctttaaaa      360
actataaaca ttgtaggaga attatagcca gtcttcagtt ataaccactc caccctcctc      420
actttctctc tctctctctc tttttttttt gctatgggat ttaatgggaa aaatatgtaa      480
aaactgtcac taa

```

```

<210> 202
<211> 283

```

<212> DNA
<213> Homo sapien

<400> 202
cctttttatc tcagtgcacac cgtccgggga cgcaggtggt ggtgactcaa ggctagcctc 60
aaagggcagc cccacctcct catcctggac cacagagacc acctgcttgg cgcgccgtcg 120
cttttccgag aggggtggctg actccggggg gctggggctg gggctgccgc ccccgccgct 180
gttgctgtac tcctcgcccc agtcgatggg ggctgccctc ggacagcagg tgcaggttgg 240
gggcaactgtt acgcaagacc atgctgcccc gagaggtaga tct 283

<210> 203
<211> 713
<212> DNA
<213> Homo sapien

<400> 203
ctgcttttgc gcaaggtgcc actggacgag cgcctcgtct tctcggggaa cctcttccag 60
caccaggagg acagcaagaa gtggagaaac cgttccagcc tcgtgcccc caactacggg 120
ctggtgctct acgaaaacaa agcggcctat gagcggcagg tcccaccacg agccgtcatc 180
aacagtgcag gctacaaaat cctcacgtcc gtggaccaat acctggagct cattggcaac 240
tccttaccag ggaccacggc aaagtcggggc agtgccccca tcctcaagtgc cccacacag 300
ttcccgctca tcctctggca tccttatgcy cgtcactact acttctgcat gatgacagaa 360
gccgagcagg acaagtggca ggctgtgctg caggactgca tccggcactg caacaatgga 420
atccctgagg actccaaggt agagggccct gcgttcacag atgccatccg catgtaccga 480
cagtccaagg agctgtacgg cacctgggag atgctgtgtg ggaacgagggt gcagatcctg 540
agcaacctgg tgatggagga gctgggccct gagctgaagg cagagctcgg cccgcggctg 600
aaggggaaac ccgcaggagc ggcaccgcag gtggatccag atcttcggac gccgtgtacc 660
acatggtgta cgagcaggcc aaaggcgcgc cttcgaagga gggggctgtc caa 713

<210> 204
<211> 275
<212> DNA
<213> Homo sapien

<400> 204
gtagacaagt acagcagatc cagacaccag atctagctag gctaaatgta cagtatctaa 60
cttgatctga actgaacctg tattccttga tgatgcctaa aactacatcc atagaattct 120
ggatgaacctg taatacagtt ctgaaagtac agttttatat aataagatgc tgatctcttt 180
attctttcaa gtaagagtgc tagagaacaa attgtgttac ttgccttgagg atttattgaa 240
cgtctggaaa atgctgtcct cctagatcca aacag 275

<210> 205
<211> 694
<212> DNA
<213> Homo sapien

<400> 205
ctgttcctgt acattttaact gaaaaaaaaaag taacttaaaa taatataaaa atagcactca 60
tgtatgtcct acagttatag gtgaaatttg atattgtttg tcttacatag catacctata 120
gacagcttaa gtaaagtgc tgttaagagg gttatgctta ttgatgaact cttgtagtgtg 180
cttaccagct ctgttagtat agttaaattg atctcagtag cttcaagtat ttataaaatg 240
gttgaagtcc aaatacatgt gataattaca atacactttg aattaatgga ggggtggagg 300
ctagttgaaa tgcattttat ttacccaagg agtatgttaa aatgatagtt ataaatgttg 360
gaagtttaaa gcaagatact cagtttagtt ctttacaaat cataagaaga acaaaattag 420

```

atgttgacat tgctatntta ggctgtgtgt tttccatatg cttcttgctt tccctgtcac 480
aggtgggtggc agcaatattg gtgtgattga ggttatgctg gcaccactcg cacacaggcg 540
cacaatgggtg ttagctgggc agaaagagtg gcattctctg ctaccgggct gggggcgacc 600
tttaccatag gatgaagtaa ccttgcatte ggctgcaagg tgtactgtac cgtacacagg 660
tgctgggtcg atggccactt tctgcttttc tttc 694

```

```

<210> 206
<211> 704
<212> DNA
<213> Homo sapien

```

```

<220>
<221> misc_feature
<222> (1)...(704)
<223> n = A,T,C or G

```

```

<400> 206
tttttttttg gnaaaaacag ggtttcatca tgtttgccag gctagtctca aactgctgac 60
ctcaggggat ttgcccgcct caccacaattc aactttcgta agtcagtatt taccatctaa 120
ctcagtgtcc caaaatttta aatttccttg cactttacag caaaaatata tattggggct 180
ctactgaagc aatatatata tgtcaaaact aaaaatcaga aaagcaaaag ggtccattca 240
acatatagca gcttatattt aaatatgtac aggtatgtat gttttcacag ttagatcttt 300
aaaaaaattt atatttgata tgttcaaaaa tacttctatt ggctataaat aatattttta 360
aagctcaact gatcaaaatg cattccaaga acatatcaaa ttaaataaat cttctacgtc 420
tttaaaaaca gataattgaa gtcagtaaag cttgaggttt gtgttaagtg tattctgtca 480
gtccctacta ctagggaagg cagaatcttc taaatacgat acgaaagaaa ctcccaaagc 540
ttggaaggaa tcggcagctc ctgaactttt tggggggggc atccctcttc gggattgaca 600
tgcgacataa atgttgcaag ctaaggggacc ccccccgggg gagtgggccc caaaaaaac 660
cacaccttcc ccgtcaatgg tggccccccc accaacctta aaaa 704

```

```

<210> 207
<211> 225
<212> DNA
<213> Homo sapien

```

```

<400> 207
ccattttaac tgtactgcca atagaattct ggaattgtgg aaaattgtat cattgaagtt 60
cagtaggatg tgtggcttaa aaatttatca ggaccacaaa aaagaaaaca aaaatatttg 120
gtactgaggt tcattgccag ggcaggaggt atttccagaa aatactcatg cctgtgttct 180
gttctctgct ttcccaaata ctgcatgtga ctttcctaag cggca 225

```

```

<210> 208
<211> 678
<212> DNA
<213> Homo sapien

```

```

<220>
<221> misc_feature
<222> (1)...(678)
<223> n = A,T,C or G

```

```

<400> 208
cctatatcta tcaaaaaaaaa tccagttcct aactaataat ctccccaaaa gaaagcacca 60
ggaccagatg atataaatgg caaatttttt caatcattta aggacaaaat aataccaatt 120

```

```
<210> 209
<211> 720
<212> DNA
<213> Homo sapien
```

<400> 209

```
<210> 210
<211> 277
<212> DNA
<213> Homo sapien
```

<400> 210

```
<210> 211
<211> 715
<212> DNA
<213> Homo sapien
```

<400> 211

qtqgtagaaa tactaatttt gcaattacag aaaaaaaciaa atgccattca catggttyct 60

```

aacaaaaagt gtctgaccac cccaccccc caccctcaa aaagccctta aataaagagg 120
aagatcaaaa gaaaacaaaa taattcccga gtttcacctc atacatacaa tatagcacag 180
gaagtggcaa agtttaaaat aatgccttta ctgttaggac tagtatgctg tcaaaagcca 240
caatcctttt gttttagtga gttgattttc aatagaaaaa taaaaatgaa catgtgttta 300
agttccaaca tggattgagc acctctgaat ttagtatcaa atgattaatt ttatttttca 360
gatgtcaaat cttagtataa aattttccat tattttaaac ttcacttgaa tctttaaaaa 420
agctgtctaa attgtactat atgagttcag tttaatcttc tgtaaaatgc taacaaattg 480
aactgtcagc agtcttttaa aaaaaaatgg gggctgggtt atttctagaa gaactctcat 540
taagctttga aaatcagaaa tcagagacaa ataacttcag atatagacta gctccacaag 600
caaatttata caattatctg taacagtcta tacatatatg tgtatatata tataccgtaa 660
ccactttcat aggtaaaaaa tattaacttc atgtcacact atgatcagaa gtata 715

```

<210> 212

<211> 717

<212> DNA

<213> Homo sapien

<400> 212

```

agcctcccc aatgccttaa aaggtcacag tagatctcag ctctgaacag aaactcaact 60
gaaactcttc ccacaaccca gcagtagata tattaataacc tacaattttc agggatacaa 120
ccaatattta attcttttga gggttttgtg ttaatacaaa ggacacaaac acacgtataa 180
aatgacgatg tcaatactga ttaaacagaa caacaaaata agaagctcaa attatcatca 240
gctatttgtg atactctgaa taacaataat gcacttgatt ctgaaagaat gatttagagt 300
cctactctga aaatctaatt gtcttgatgt ggcgaagtga gaagaaagga tgatttttct 360
aatgaaaagc atgtatacgg gttagccctt ggcgagattct gtcaaaaccc tgaattttgc 420
attagctgtt ttaccaccca aacgttttta cccgaggatg tgcagcaatg ggaactctca 480
tacactgctt gtgggaatat aaatcagtat aaccactttg gaaaaccatt taacattgtc 540
aactacagct ctacacacaa gtgctataac caccatttcc actccagggt atacacccta 600
aaaatatgaa gtgcccattg ctacccaaaa ggccgcctaa aagggaatgct tttgagaagg 660
gttaaccttg ttaattagtg gcaaaactgg gaaaacaacc cccaaatggc cccatcc 717

```

<210> 213

<211> 599

<212> DNA

<213> Homo sapien

<400> 213

```

cctgttttgg cgaggcagga gggaaagcgg atgggagtg tggtaggcc aagggtagtt 60
caaagcgatt cagcaggatg atgaccacag gagtgtgga gccgggcctt tcagcccccg 120
tgtggatgat gaccggccat ccaggacatg cgagggtctg ggacagtgga cagccagtgc 180
cacacaagga aggaccgatt aaatgacaca gttaaaggaa tttggcctag ggagtgcagg 240
ccagaaaagg ttggtctttt tatatatgta acattggaaa aaaggaacat ctctgtttcc 300
ctgtattaag ttttgacttt agctcagcaa atgcagtgtt tgtggcagta aatatactct 360
gataacaatg ttctttccca ggaatttaga gttttatgat ggttattgaa aatgtttaca 420
tgacaggctg tcaataatat tttttgctc taaaaataaa acatacataa agtgtaggga 480
ttttaagtat gcaactcact gaacttttca taccgtaata caccacccta gtaaccctcc 540
cccagttcaa gatgtagact gtttccaata acccctcatc ctgttcccta atagcccc 599

```

<210> 214

<211> 789

<212> DNA

<213> Homo sapien

<400> 214

ccttatgaca	aaccttgcta	tgccaaggat	atgcttcact	atcttcatct	atcaaaacac	60
tatgcatcat	agatatctaa	ttttttcatt	tcttgcatga	agtctttcct	gatttccctc	120
tgctgaaatt	tctctcttca	aatgatgtgt	ttccatagta	ctttgtccct	tttcaaagat	180
atatctcaca	tcgcatatct	taccacagtt	agtttcattt	cttaactctc	acactagatt	240
acaaagtcaa	tatagacaaa	gaaatgttca	accttatata	acctcctctg	cctatgctgg	300
taaattgcac	ctactatgtg	ttcaataaga	gcttgtcttt	ttcaatatac	aaaactttgt	360
aaagattaaa	gaccttgtag	aaagtcaaga	ggaagatagc	aatttcactt	ctaagaactt	420
accctaagga	aacattcatg	aagagataca	aggggttatg	tgcatggatg	ttcattatca	480
tattattctt	cattatgaag	attatgatgg	taataatgaa	aatgattatc	ttgtattggg	540
ccttatttga	agtcaagcat	tgagaatgta	ctttatctgc	attatctcac	tgagttctcg	600
tagcagccct	ataagggtaca	gactgttatc	taagcttaaa	aaaataaagt	taatgtccaa	660
ggtcaaacaa	ctagtaaaaag	aagggggcta	ggaaatttgg	aacccccaaa	ggggcaacct	720
ctcaagggct	atgaatcctt	accattatta	taaggaagct	tggcccatgg	tggcccaaaa	780
aaaaccggg						789

<210> 215

<211> 765

<212> DNA

<213> Homo sapien

<400> 215

ggatgtctga	gcaggagaga	gaccatgtga	aggatggact	gaatggagac	ttgtatcaaa	60
gagtcctgag	atcaaagact	tgtattagag	aggggtgttg	tagtaatcta	gtcaggggtat	120
gagaaatggg	ttgtattaga	gtgtcaggag	tagtcgtggc	aaaaatatat	agatcaggat	180
gagggatggg	cctcatctca	cacctgact	ccagtcaatg	gcagtggctc	cctggagtag	240
actactatag	gaaggatttt	gtaaaagttt	gtctggcctc	agtggagggt	gaggtagggg	300
aggagttcta	tgaacagtta	gtggtgtctg	ccatggttga	aacaatggag	aagggggaca	360
ccttttctgt	gcagatgttg	cttctggtag	atataatcca	caatgtaatg	ggagaagtac	420
taagaatcag	taaattatgg	aggggtgtaa	agactactga	tatttaagcc	tgcggaaccg	480
acttagagaa	atgatagtta	aaggagaaat	atccagcaaa	caaagatatg	acattgaagt	540
ttgggactgc	gatttagtacc	agagatttgg	attggagggtg	atttgtagat	aatggatagg	600
tgattttact	cttgcaattt	ggattgaggg	gtgggggaaa	ccagaaaggg	gctggggggg	660
aaattagtag	aagggtcacct	tgaattcatt	gtggtccata	tcaatgctga	aactgattgg	720
ggaacttttt	actcttgagt	ccctttgtaa	gggaacccca	gaaag		765

<210> 216

<211> 780

<212> DNA

<213> Homo sapien

<400> 216

cctttttctg	tggcaaatgg	aggctttttca	ctgcctgtag	agacaatata	gtaagcatag	60
ttaaggggtg	ggtcagaaca	tgtaagata	acttactgta	tatgtattcc	cttgtatttt	120
gttaaagctg	gaacatttga	tattttttcca	tttatttatg	aaaaaatatg	aacctatttt	180
cattttgtaca	aggtaattgt	tttttaaaagc	aagtcacctt	aggggtggctt	taattgtata	240
agtcaagcac	atgtaataaa	ttcaaaacct	gcagttaaca	ggatattaga	catcaatcct	300
ggtaaccaa	tattaaagat	tctcttttaa	aaagactgaa	catgtttaca	ggtttgaatt	360
aggctaaaag	gtcttgacgt	ggcttttcat	ggcccttcaa	attggaatgg	aactactgta	420
ctttgcccatt	tttctataaa	tcagtacttt	ttttttaatt	ttgatataca	ttgtgtgaaa	480
aaagaaaatg	gctaataaac	tgtattaaat	cttaaacaat	gtataaaagat	tgcaacttagc	540
cagttcaaag	tgtatactta	ttcataatga	attataacag	ttatatattct	gtgttttctt	600
gtaaatgttt	cttttccctt	aaatacagat	aattcatttg	tattgcttat	tttattatga	660
gctacaacaa	aaggacttca	ggaacaagta	atgtattagt	atgggttcaag	attgttgata	720
ggaactgtct	caaaaggatg	gtgggttattt	taaataataa	tagctaattgg	gggtggtaaa	780

<400> 217

```
<210> 218
<211> 817
<212> DNA
<213> Homo sapien
```

<400> 218

```
<210> 219
<211> 661
<212> DNA
<213> Homo sapien
```

<400> 219

ggatgctgag	gcaggaggat	tgagtccctgg	agtttcagga	tacagtgagc	tatgatcatg	60
ccattgcact	ccagcctggg	caacagagca	agattctgtc	tctaagaaaa	ggaaaaagaa	120
aatgaataga	tagtgggtatt	agatgttaat	gacatcagtt	gtttttattc	tttattcttt	180
cttagaaaaca	gattagtttt	ctcgaattaa	agaactacca	tttttctttt	ttctacaact	240
ttcaagagct	ggtgaagaaa	tgatgttttag	atttaataga	tatagtagca	gtcatatatt	300

```

aatagaatag aaactgagac tctaggaaaa agatagacat gagataagga gtaggcatgg 360
tagacatttc tagattatct atgaaaatgt tgtagaattc attttttttt ttggtctgac 420
ctttggcaat ggtgctgagg aagggaaaagc cagcccatca ggcaaggctc tgttttctgc 480
attttatccc gtttgattct tctcgtagg attggagcaa ataatttcaa tatgttcttc 540
gctgggttta tcatagtac ctttcattta aagggaacttt taacaattga cttaaagaac 600
actgagatgt gatattttat tgggatttga aagttgccat tgggttttac cttccttaat 660
t 661

```

```

<210> 220
<211> 792
<212> DNA
<213> Homo sapien

```

```

<220>
<221> misc_feature
<222> (1)...(792)
<223> n = A,T,C or G

```

```

<400> 220
cctcttttta ttcctacaaa taattttcaa gtacacacaa ttgggtaaac aaagaaacaa 60
agccaccaag aatgaaaatc agtaggaata acgaacaaga ctcacagatg tcaaacaagt 120
ctgtgggtct tgcagacttc agatgttgga attattagtc gtggcaagng nncaaacat 180
tagctattac cattatgttt accaactagt gaagtgaact atgagaggat atattaacca 240
cagaagttaa tagaagaata gactcctgaa aatatctgga tgctacaaac taaaatatag 300
tatataatcc ttcatagagt gtcagtgact tcatatttat aattacattt ttgtatatta 360
gcagtgttct agttcttact gccttatctt taagctgann nnaaataaaa ttatatattg 420
ggattcaaaa acacatagct aatgattact atgtggcagt gttacattac tttatcacat 480
atcattaaca taatctgcat gtgttcaaaag agatcttcat acttctttgt agctccact 540
tctttgtcgt cttttagctt cccacaacat ctagaacagc acaaccgtat atggagaaaa 600
ctcagtctag tattcgttga atgactaatg gaaaatttag ttnataaaca gaactttctt 660
cattgnacaa attatcttgc agaagaataa tggccttagt ttaaaattat catatttacc 720
catntcncca ngttatttta tctcttttgg ctaanaattt tgaaaacggt accttttacc 780
ctttggcatt tt 792

```

```

<210> 221
<211> 759
<212> DNA
<213> Homo sapien

```

```

<220>
<221> misc_feature
<222> (1)...(759)
<223> n = A,T,C or G

```

```

<400> 221
cttttctgct gctccgggag gtggagtggc ctggcagagg gcacatggct gccacctgct 60
gcaaggaaaa ttctcagtga agactcctca gtatgaagga gataagcctg cacaatcagt 120
cactgataga tgcttagtgg aaaaacttcc aattcccatt tacagctctc agagctagga 180
ttaaaaaactc ctggtcataa actcatgtga tgagaagtta tagcacgccc tcattttcta 240
catanccact tgcatttatg gttggctttt gaacttgcta gaagggaag aagtgcaa 300
gtgtcctcct tagagctact ctctccctt tgggtgggtt ccagtttggt cattgtccag 360
atggcccagg agctgacgat caaagggaag aagtcatgtt tgtcatgaga atgctttgct 420
gcatcaggat tcagtgaagc tggtcacgc ctggagccca tgcagcctca agaggcagga 480
tggagctcag aaaccatcac tgaggttaga aagtgagcac caaagttgag ggaagccac 540

```

```

aggagtgagc cgaagtgctc cctttggatt tccaaagtgg gtgctgctgc ttcttccatc 600
agccttgctt ctgaccccaa tgcgttcctg gtgccttctt cttggcattt tgctgtcggg 660
ggcccaagga aaaaaattcc tgcattggcag tggtgaaaaa agatggctgc ctgctgaaac 720
ctgatttggc ctgggtaagc cttttggagc cccgggtaaa 759

```

```

<210> 222
<211> 699
<212> DNA
<213> Homo sapien

```

```

<220>
<221> misc_feature
<222> (1) ... (699)
<223> n = A,T,C or G

```

```

<400> 222
ccttntnaag agttggcatt aattcttcac taaatgtagg agtagaattt atcaggtaag 60
ccacactgac ctctggnctt nttnncgccc gatgattttt aattagttga atccctttac 120
ttgttatata tgtattcata tattctgttc cttcttggat ttacttttat gattggtgcc 180
tattgaggta tttatttcta gtttgtggta cttcatgtgt ttaggttttc tagacagtgg 240
acatagaaga ttcaagaagc taaatgtagg agaatgtnta atgtaggana ntgaggcnac 300
natatcatca atgaatgact tgaagtttcc tctgttgtaa agaatgatat taccataact 360
gccatagnta atattgatgg tgtaagtcaa ataanaaggc aggaggaaaag ggacatccat 420
cactgaacca canatcagag nctcattgaa gcctttgaga agaatccaca aaattttaca 480
ggataattca tttctgcgga tcaccacnag aagagaaact ggttaaacag acagggtattc 540
cagagtccaa aaatttacat ttggtttcng aaccaaagac ctcagctccc aggccacagc 600
aaaagggggc ttatgaattc cctggcaccc agncccaaga cccaanaacc tcactttgat 660
tggtttnggg cttgggaaac caaaaaacca atgggtgggc 699

```

```

<210> 223
<211> 598
<212> DNA
<213> Homo sapien

```

```

<400> 223
aaaaagagaa agtttcagat ttgccattca aggccttattt atatatatgt gtgtgtatat 60
aaatacatgc acacacttgc atacatatat atttttggct gggggagtggt gagttttgcc 120
tttctaaggg agggaccgcg caggctcctt tgttctgtat tctggcggag atgggtcctg 180
gccttggtgc actggcctat ccttaaagat catctcccat cctcccagc gccatctgtg 240
tgcagcaacc agaaagggat gaacttggcc ctcttgcggt cctggacaag gtctcttctt 300
taccctttct gttgccagtc agcaacctgt aactcacatt ctcttcccag tgaatccctg 360
ggagcgcttg accctgggtg gctgttcagc ttcctgctgc tggggccagc aatttttgag 420
gatttatctt taggccaggc ttgcctccgt acttatccct gctctcccat ttctctcttg 480
tttgagagag aatgaggaag caaagagtga gaaagaatag gggctgaaga cgccactccc 540
agatggctct ttctatctg ctcttctgtt gaaacacacg tgctgtgggc ctcaggcg 598

```

```

<210> 224
<211> 501
<212> DNA
<213> Homo sapien

```

```

<220>
<221> misc_feature
<222> (1) ... (501)

```

<223> n = A,T,C or G

<400> 224

aaacctttat	gatgacttcc	ttatgaatta	ctgaacgaac	actggaatgg	gactcaggtg	60
tcttgaggac	atctctcaac	tctggcctta	gttccccctc	tgtaaaatta	gggtgccaac	120
taaatgatct	acaaggtccc	ttccagcgcc	gccatttctgt	aattacatca	tgtgtaactg	180
tattaaacat	acacaagtga	ctgccaggca	tgggaatgta	acttccgagt	aaatgctttg	240
gtttgttcag	aatacactat	gaacttcttt	ccaaagacgg	gttgtggtaa	atagtggata	300
ttttgattat	aagaaataga	gtttccttga	agcttttagct	ggagatacag	caatagtgtg	360
gtgttcctac	aaatatcaca	gtgtattcaa	acatatTTTT	ctatcaaaaa	tcattttttgt	420
aaaagctgtg	tgtttttatc	caacttgtga	taataaatgt	tctttatttt	agaacaaana	480
aaaaaaaaaa	aaaaaaaaaa	a				501

<210> 225

<211> 295

<212> DNA

<213> Homo sapien

<400> 225

cctgtatagg	gctcgtttcc	ccacacatgc	ctattttctga	agaggcttct	gtcttatttg	60
aaggccagcc	cacacccagc	tactttaaca	ccagggtttat	ggaaaatgtc	aggaaaaaaaa	120
aaaaaaaaaa	cacatgcact	cacacaatac	ccaaacatca	raattagaag	ggcataaaac	180
agggggcctt	ataggctgaa	aaatatctta	ratttcaraa	cagaatacca	atcaaatatt	240
gaaaattcct	ttgttcaaaa	cacaaagatg	ttttgttttt	aatggggagtt	ttttt	295

<210> 226

<211> 372

<212> DNA

<213> Homo sapien

<400> 226

agattcctgg	cttagagcat	gcgagcattg	aaggaccaat	agcaaactta	tcagtacttg	60
gaacagaaga	acttcggcaa	cgagaacact	atctcaagca	gaagagagat	aagttgatgt	120
ccatgagaaa	ggatatgagg	actaaacaga	tacaaaatat	ggagcagaaa	ggaaaaccca	180
ctggggaggt	agaggaaatg	acagagaaac	cagaaatgac	agcagaggag	aagcaaact	240
tactaaagag	gagattgctt	gcagagaaac	tcaaagaaga	agttattaat	aagtaataat	300
taagaacaat	ttaacaaaat	ggaagttcaa	attgtcttaa	aaataaatta	tttagtcctg	360
atgaaatgaa	at					372

<210> 227

<211> 599

<212> DNA

<213> Homo sapien

<400> 227

ggcccccgtc	gcgggagccg	cttcgggcct	tctgggcatg	tctgcatat	ggctccaggt	60
ttgtttttct	ccccggcact	ctgacgggga	gggtctcccg	catctcctgg	catccgggta	120
gaggacgcgg	aggatgctga	gctgctggcg	cactgcagca	caactagaga	tgtacggatg	180
cccccatctt	gatcttacag	aatcagaggt	acagccgcga	gaaagagtca	agaacagaca	240
gagtcgcttg	aggactcagg	agggtgtttg	ctgcgttgac	aacagactac	accctcacag	300
tttgctctgc	tcttccaaca	ccagtgggaag	atgatcacat	cccagggatc	agtgtcgttt	360
agggatgtga	ctgtgggctt	cactcaagag	gagtggcagc	atctggaccc	tgctcagagg	420
accctgtaca	gggatgtgat	gctggagaac	tacagccacc	ttgtctcagt	aggggtattgc	480
attcctaaac	cagaagtgat	tctcaagttg	gagaaaggcg	aggagccatg	gatattagag	540

gaaaaatttc caagccagag tcattctggaa ttaattaata ccagtagaaa ctattcaat 599

<210> 228
<211> 343
<212> DNA
<213> Homo sapien

<400> 228
aaagtaaatt gtatgaaaaa ttcattttctt caattgcatt agccacattt tgagtattca 60
tgtggctggt agattctgta ttagcacaaa gatatggaac atttccatca ccacagaaag 120
ttctgttgga cagcactgca ttagaatatt ttcatactgc tcttctctcaa ttaatttttg 180
ttgttaattgt tgatgtcttc attggatggg tcataatgtt ccatgaaacc gctcaagtac 240
acaattgtat gttcttttga tcccttacca caaatatctc gctctgctca tttcttttgc 300
agcttcctat aaagtttgtc ttctcaaaa aaaaaaaaaa aaa 343

<210> 229
<211> 417
<212> DNA
<213> Homo sapien

<400> 229
ctcaagctgc agtccaccgg gtatggttct ggatggttcc cccaagggag caggtatgta 60
ggaggtgaag aaaactgaga tttcaagtat gggagagttt ttactatctc cattcctgga 120
ttaaaagtgc tgaaaaagtc cacagttaaa cattccttta ttcacctat ggctcccaag 180
aaaagcattc ttcctctgga gtactgggtg actaagggga caatacacca aatttggtga 240
gtttacaatc aagcttacta aggttggtgact tcttatcag tttggcagag tcccagggca 300
gaataatcat ccacttacag gtctctgttt cctctccctc cgcagcagtg gagagcatcc 360
cagtgtttgg ggcactgtgt tcctcttcgt ccctgcacca gacctggaa gccttgg 417

<210> 230
<211> 462
<212> DNA
<213> Homo sapien

<400> 230
gaaataccag aagagaaagt ttcattgtgc aaatctaact tcatggcctc gctggctgta 60
ttccttatat gatgctgaga ccttaattgga cagaatcaag aaacagctac gtgaatggga 120
cgaaaatcta aaagatgatt ctcttccttc aaatccaata gatttttctt acagagtagc 180
tgcttgctct cctattgatg atgtattgag aattcagctc cttaaaattg gcagtgtctat 240
ccagcgactt cgctgtgaat tagacattat gaataaatgt acttcccttt gctgtaaaca 300
atgtcaagaa acagaaataa caaccaaaaa tgaaatattc agtttatcct tatgtgggcc 360
gatggcagct tatgtgaatc ctcattggata tgtgcatgag acacttactg tgtataaggc 420
ttgcaacttg aatctgatag gccggtctc tacagaacac ag 462

<210> 231
<211> 328
<212> DNA
<213> Homo sapien

<400> 231
ctgtgggttt tcttaaagc ccctcatctg gttgaagccc tagtgtttct ttctcacatc 60
agaggcaaag gcattggggg gggctctggt tggacaataa atttccctctg gtttggacca 120
agaaaaacag agttctttga ccgctaact atattgtaaaa agaaagtgtg taaaaacaag 180
agttaaaatg cttctaacag tgtggctatc actgcacagg acactggaat tggcattcgg 240

ggttgtgtct gtccatgtgg tttcgttgta tgtcatgtgc tctcagctca gacagagaca 300
tccaattgac ttctgacttg gggcattt 328

<210> 232
<211> 595
<212> DNA
<213> Homo sapien

<400> 232
cgccaatttt agcaaataag agattgtaaa agaagcagat tgaatgaaga attttttagct 60
gtgcagatag gtgatgttgg gatggaaaat gctaataaac taccctttct tttatcaagt 120
aattaaaata aatctacata aagaaccaa aaggctgttt tataaaagt aaatatccag 180
tatttcagag ggccaggcaa gagcacttca gatgaggcag tcaaaatcat tttttccag 240
tgaggataga ccacaagtgg gtggtgagac cattgaaagc ctttatcaac tgaagagtcc 300
atttaacagc ataatttgtg ggaagactgg aatagggctg aataaatgtg tttgaatctc 360
taattttata ctttcttttc ctgaggaact tgatttttct gtccctggat cgccttgtca 420
taattgggtc tgttcctttt actaccactc ttgagtccat atatgaaatc attaaagttg 480
gatgatcagt tttttataaa aatatatatt tttgtccaag aaaaaaaaaa gcatacatat 540
gtgattatgg ctaaatacaa ggtaactgga atgtatatac ttttgctaatt gttcc 595

<210> 233
<211> 600
<212> DNA
<213> Homo sapien

<400> 233
atgaaggtaa actctaaaat cttcataggt caacaaagaa aattttatcct tcacacttat 60
ttctagaaag cagcagggct tatttcctag attgcttaca atgaagctag aatatctgcg 120
ataactgtag agtttcaaaa aggatcccta gggctacttc tacgttctcc ttaccagttg 180
agcactctcc ataatttcca gacgggtcat gggggagaat gatagaaatg agcgtgggaa 240
gaaagacaat gaaattagaa atgggtgaga cacatggtgg tagaatgcta agagcaggggaa 300
tcaggacaat caaccagtg tctaggaagg gtcaagtcac cagtgtcatc tgctgaccaa 360
tgttaggaag aaataaactc aaaggaaaaca ccacattttt ccaattaaac tcaaacttat 420
tgacttgtgg tggttctttg atgttgtggg gactgctata acagaaaacca attggatttt 480
caagggcaag aaactttgcc actgaataag atgatgtcat ccttctctgat aacaaatagg 540
aatgggtggc cagctctaaa cagcgtggac tgagggagtt gcttttctac aatattactt 600

<210> 234
<211> 500
<212> DNA
<213> Homo sapien

<400> 234
aaatttcctaa ttcttttact atcttctcaa cttttcccaa agataaaata aatttcacat 60
aatttcattg aggggaaatg gtagttgtaa aaaactacct caagtagcaa tcaccgctgg 120
cagtgttttc tcactttctg ttctgcaatt gcaatcacac ttccaaaaag aaaagcaaat 180
gtttgctaaa ccatagacag acaacctctt tgtgactggc attataaggc ttataatgaa 240
aacttatcaa atataaaagg tgctccctct tgaaaatgtg tattttatatt gaagttttga 300
gtaagaggtg agtgtttggc aattttcaac actcccctca aaaatctccc aaagttgcaa 360
aaaagtcagt ttagtaaaat tccaagcact taaatgcttc attgagggcc agttgatata 420
cgcaatgcac taatgtgtaa aaattaaccg aatgcaacta ttttataatg gagagctctt 480
accttttctc tccagttttt 500

<210> 235

<211> 159
 <212> DNA
 <213> Homo sapien

<400> 235
 aaaattttaca gataaaggca gttcaatact gccactgaga agtacatctc ttaacatata 60
 caacttttcag gccacagttt tgaaggctctg aagtattaag ttggtttgat gaattagtcg 120
 gttggcactt acgaacacat ttattgcctt gccatcttt 159

<210> 236
 <211> 254
 <212> DNA
 <213> Homo sapien

<400> 236
 aaataagtga ataagcgata tttattatct gcaaggtttt tttgtgtgtg tttttgtttt 60
 tatttttcaat atgcaagtta ggcttaattt ttttatctaa tgatcatcat gaaatgaata 120
 agagggctta agaatttgkc catttgcatt cggaaaagaa tgaccagcaa aaggtttact 180
 aatacctctc cctttgggga tttaatgtct ggtgctgccg cctgagtytc aagaattaaa 240
 gctgcaagag gact 254

<210> 237
 <211> 591
 <212> DNA
 <213> Homo sapien

<220>
 <221> misc_feature
 <222> (1)...(591)
 <223> n = A,T,C or G

<400> 237
 tttttttttt tttttttttt tttttttcta atttttactt tttctcaagt ttaatgtara 60
 catacaaraa aacatcaagc aatggtttatt gkgcaattcc aatcattatt tgcaraatct 120
 tggtttaaag tcagtyttta tagccatttc aactgcttgg tttaaacaaa aagcaacaat 180
 ctggttatyt acctataaat ttcatgggat ttytttaaac actgaagtac taaaagcact 240
 gatgatttgt attataattt ttaaaatatt taaaacctac acagatttca taratcattc 300
 cttttataaa ataatcaaaa taatttgatt atytggaata aaaaattcctt gaaacaragc 360
 cctttccagg tatyttcaat ctctgtaaaa ccccaaacc ccaacagagt aratgatgaa 420
 ataaggattt ctcagttgcc caagactgtc tgaaatttaa ggttgaaaaa tggactggcg 480
 tttttcatgt ttctgngaa ttcanagctt acagggtggc tcaaaactca aatctctggg 540
 atggctttac atggctttca ctttgatttg tttcattttc atttgcttct t 591

<210> 238
 <211> 252
 <212> DNA
 <213> Homo sapien

<400> 238
 aaatggcttt tgccacatac atagatcttc atgatgtgtg agtgtaattc catgtggata 60
 tcagttacca aacattacaa aaaattttat ggccaaaat gaccaacgaa attgttataa 120
 tagaatttat ccaattttga tctttttata ttcttctacc acacctggaa acagaccaat 180
 agacattttg gggttttata ataggaattt gtataaagca ttactctttt tcaataaatt 240
 gttttttaat tt 252

<400> 239

$\langle 210 \rangle$ 240

<212> DNA

<213> Homo sapien

$\langle 400 \rangle$ 240

<210> 241

<211> 400

<212> DNA

<213> Homo sapien

<400> 241

<210> 242

<211> 75

<212> DNA

<213> Homo sapien

<400> 242

<210> 243

<211> 192

<212> DNA

<213> Homo sapien

<400> 243

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gctccacatt tgtagcgaac actttgactc caaagagaag gaggaagaca aagacaagaa      60
ggaaaagaaa gacaaggaca agaaggaagc ccctgctgac atgggagcac atcagggagt      120
ggctgttctg gggattgccc ttattgctat gggggaggag attggtgcag agatggcatt      180
acgaaccttt gg                                     192

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```

<210> 244
<211> 616
<212> DNA
<213> Homo sapien

```

```

<400> 244
aattttatag caatatactg accatttctaa aaataacaaa atacatgttg ctctcaacta      60
catagttaaa aaaggtagta aattctctta cccaaaatag aggaggggtg ggctagttag      120
ctgctcaaac atttgtaaca aataaaaaatg tatctatata catataatga tcatgttttc      180
atagcctaaa atcaccatac aaaatctaata aataaaattg tgtcgtgttc aggagttggg      240
aagccaacac attaaattaa caaagtattt ttggtatatg taaataatgg gatagaatct      300
ctcgaatcag gattgtccca gaagttctaa ggcagatgtc aatgacatgc acattgtcca      360
tggttcagtaa ttttcaaaga ctagaataaa ctatgtaaac tattcaatac aattcaatat      420
tacttaactg ctaaaaagta cttcaagatc ttgcactgcc ttgagttagt ataatcaaat      480
tagtaattgg aaaatagctg taatagcagg cactgaagaa ttctgacaaa taccaaataa      540
ctgtttgttt ttaccaaata aactggtaag atgatatcac aaagggtttt aagttatttt      600
gctatacaag gttttt                                     616

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<210> 245
<211> 165
<212> DNA
<213> Homo sapien

```

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<400> 245
ttggaacagt ggattaaaat ccagaagggg aggggtcatg aagaagaaac caggggagta      60
atttcttacc aaacattacc aagaaatatg ccaagtcaca gagcccagat tatggccgcg      120
taccctgaag gttatagaac actccaaga aacagcaaga caagg                                     165

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```

<210> 246
<211> 229
<212> DNA
<213> Homo sapien

```

```

<400> 246
tgtactggat cctccaggt gggggcgact ctcacctgac tattacaata gcctcctaag      60
tggtttccct acttgcaacc ttgccgatat aatatctatc ctccacacag caggcagggc      120
gatcctttta gaatagaagt tagatcatga aaatgctctg ctctgatccc tgcaaaagct      180
cgccacctcc ttacagtcac cgctgaactc gtagcagagg ttcaggagg                                     229

```

```

<210> 247
<211> 338
<212> DNA
<213> Homo sapien

```

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<220>
<221> misc_feature
<222> (1)...(338)
<223> n = A,T,C or G

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<400> 247

<211> 177

<213> Homo sapien

<400> 248

<211> 263

<213> Homo sapien

<400> 249

<211> 333

<213> Homo sapien

<400> 250

<211> 384

<213> Homo sapien

<400> 251

agacaccaac tcgtttctag agggctaaga actgcacttt aagaaagggc ggggaggtga 360
 agggacccga gcaagaactt tcag 384

<210> 252
 <211> 211
 <212> DNA
 <213> Homo sapien

<400> 252
 aaagcagtct gaaaatggga catctgtaga gaaattcatt tccttcttct cctccggatg 60
 tggaatggaa gctttgaggg aaggaaaagt aggaaaagag cgggatggga tgggatggga 120
 tgggatggga tgggatagga agagaggctg gggaatgggc agagaagggg gtgctgagtg 180
 tgtgtgaga tagagcaaga tcacaagaag g 211

<210> 253
 <211> 135
 <212> DNA
 <213> Homo sapien

<400> 253
 aaaaattggt tcttgacaag ctgacttggc acttaagtgc acttttttat gaagaaaaag 60
 tacaatgaac tgcttttctt caagcaataa ttgtttccaa cttgtctggg aattgtgtgt 120
 ctggttaactg gaagg 135

<210> 254
 <211> 361
 <212> DNA
 <213> Homo sapien

<400> 254
 cctgtagccc ctgctacacg ggaggctgaa gtgggaggat cacttgaacc aatgaggggtg 60
 aggttacagt gagccagat catgccacta ctctacaggc tgggtgataa gagtgagacc 120
 ctgtatcaaaa aaaaagacaa ggaaaaaaaa aactgggccg tttgtttttg cagaatgtct 180
 ctcaattttgg acttttttggg caggaatata atacaagtga taaaaatgct tctttaacat 240
 tagaacctgt ataaaattac cattacagac cttgctatct tacttatagg taaatcactg 300
 tttaccaagg taagtctttt gggaatttcc aaaaatgaag tccatggaca gttaaaaact 360
 g 361

<210> 255
 <211> 331
 <212> DNA
 <213> Homo sapien

<400> 255
 aaaaaataaa ataatccacc aacgtgattg accttggcga gatcatgttt ctagtctata 60
 cctcagtttc cccatctgta aagtgaggat aatgtcccac cccatgtaac tgtggtgagg 120
 accaactgca aactgtgcc tgcgagtctc cttggaaaaag tgtaaggttc tacacaaatg 180
 gaaagtgatc tgatcacact cagtgtcccc agcccagcct ttcagtgcc tggccctggg 240
 gtgggggaca atactctcct caccctcttc actagtcttc atgaatagca aggaggccat 300
 aacataattt ggtctaaacc ccttcctttt t 331

<210> 256
 <211> 186
 <212> DNA

<213> Homo sapien

<220>

<221> misc_feature

<222> (1)...(186)

<223> n = A,T,C or G

<400> 256

ccttttgggcc	cttgcacttt	gacctgcaat	ggggccacac	cagccttgct	tgtgtccacc	60
tggaaggact	gagggagggt	ggcacgaacc	atgcctgggc	tcaggccggg	cccanagcac	120
ttgaccttgg	acgcatctgt	cacatcatgc	acagggacct	tgaaaggact	gcctggcact	180
tgatgg						186

<210> 257

<211> 255

<212> DNA

<213> Homo sapien

<400> 257

ctgggggtccg	tcaccgacct	ttggggaact	gggctacggg	gaccacaagc	ccaagtcttc	60
cactgcagcc	caggaggtaa	agactctgga	tggcattttc	tcagagcagg	tcgccatggg	120
ctactcacac	tccttgggtga	tagcaagaga	tgaaagttag	actgagaaa	agaagatcaa	180
gaaactgcca	gaatacaacc	cccgaaccct	ctgatgctcc	cagagactcc	tccgactcca	240
cacctctcgc	ggcag					255

<210> 258

<211> 604

<212> DNA

<213> Homo sapien

<400> 258

ctgaatttgc	aatggagttt	ggtggtgcaa	tcggtattga	ttagtttggc	atagacagat	60
gcagcagttt	agagcaaaat	cgagaaaatg	atTTTTTTTT	tcctccttga	tttcctggca	120
gaagatatct	tactttttca	gcaaaactttt	cttttaacac	taaagcagcc	tagggcaatg	180
ccagatactt	agagcttttc	tcttgattat	aagtagaaat	gggggtgtct	gggctagagg	240
tggagggtgg	atgtgctgtc	gtcacagtct	agctggcagc	aagcaaggca	aaagcagaga	300
ctgctctaga	agcgggtcca	agcagcagag	acgtcaggaa	aggcacttct	tagtaccac	360
ctctatgctt	taatagttgc	ttgttaagct	gcttcatggg	ttgagacaaa	ctaccagcac	420
ttcaaagagc	tcagttctct	gctcaactct	cttctctagt	tacattatct	tttttccttc	480
aggagactga	ggcaggaaaa	tcgcttgaac	tcaggagggtc	gaggccgcag	tgagccaaga	540
tcacaccacc	gcactccagc	ctgggccttg	caaagtgtct	ggattacagg	aatgagccac	600
cagg						604

<210> 259

<211> 429

<212> DNA

<213> Homo sapien

<400> 259

aaaaatgtct	gtatcgagat	cttccagttt	gaagtcttcc	tcctctgtgt	cttcccaagg	60
ctctgtggca	agctccactg	gttctcccg	ttocatcaga	accactgact	tccacaatcc	120
tggctatccc	aagtacctgg	gcacccccca	cctggaaactg	tacttgagtg	actcacttag	180
aaacttgaac	aaagagcggc	aattccactt	cgctgggtatc	aggtcccggc	tcaaccacat	240
gctggctatg	ctgtcaagga	gaacactctt	tactgaaaac	caccttggcc	ttcattctgg	300

caatttcagc agagttaatt tgcttgctgt tagagatgta gcactttatc cttcctatca 360
 gtaactgctc cgtgttcaga ctcttggttt cttccaggct tacagtggac atcatcagct 420
 tcctgcttt 429

<210> 260
 <211> 385
 <212> DNA
 <213> Homo sapien

<220>
 <221> misc_feature
 <222> (1)...(385)
 <223> n = A,T,C or G

<400> 260
 ctgcaacaca tgcagcacca gtctcagcct tctcctcggc agcactcccc tgcgcctct 60
 cagataacat ccccatcccc tgccatcggg agccccccagc cagcctctca gcagcaccag 120
 tcgcaaatac agtctcagac acagactcaa gtattatcgc aggtcagtat tttctgaana 180
 cgcataatggc agacggattt gcgtatacca aggagagtgg cataggaggg aaaagcatat 240
 gtggctgaaa cctgtaagtt ggtgttggtt atgcagaaat gtgtaacaga tcaaacgggtc 300
 ctctcaagtg tctattanat aggcaataag aactgcagtg tagctgagta acatctttta 360
 gctgactata aatcactttg ttttt 385

<210> 261
 <211> 230
 <212> DNA
 <213> Homo sapien

<400> 261
 ctgtactgga tccctccagg tgggggcgac tctcacctga ctattacaat agcctcctaa 60
 gtggtttccc tacttgcaac cttgcccgtta taatatctat cctccacaca gcaggcaggg 120
 cgatcccttta agaatagaag ttagatcatg aaaatgctct gctctgatcc ctgcaaaaagc 180
 tcgccacctc cttacagtca ccgctgaact cgtagcagag gttcaggagg 230

<210> 262
 <211> 198
 <212> DNA
 <213> Homo sapien

<220>
 <221> misc_feature
 <222> (1)...(198)
 <223> n = A,T,C or G

<400> 262
 atgttaagta aacatgaaat ctatataaca gaacaaaaat tcactcttat gtcaatgtca 60
 gcgtgttaat gtagatctat ttactganac agactctgta gtggcagaga gtggccttgt 120
 taagccagga cctgttctg caggctgtgg gtagaagcta ggaagtccct ggagtttcac 180
 ccagcttttc catgaatg 198

<210> 263
 <211> 157
 <212> DNA
 <213> Homo sapien

<400> 263
 aaaatatatt tctaaacaga atggggccgac tcagtcacag taactgttga tctccatagt 60
 agagcaaccc acaaagacag aactgatttt tttcccataa tcaggggtga aaaatatata 120
 acttgtttct gaacccaaaac cacaatttct gcagttt 157

<210> 264
 <211> 290
 <212> DNA
 <213> Homo sapien

<400> 264
 ctggctactc caagaccctg gcattgaggct gaggacaact tacaagggtc tcaccgaagc 60
 agtggacctt tattttgacc acctgatgtc caggggtggtg ccactccagt acaagcgtgg 120
 gggacctatc attgccgtgc aggtggagaa tgaatatggt tcctataata aagaccccg 180
 atacatgccc tacgtcaaga aggcactgga ggaccgtggc attgtggaac tgctcctgac 240
 ttcagacaac aaggatgggc tgagcaaggg gattgtccag ggagtcttgg 290

<210> 265
 <211> 234
 <212> DNA
 <213> Homo sapien

<400> 265
 aaaaaaagga aaggaaagag aggaaaagaa aataaaataa gacgatttat tgcttctcct 60
 cagcatcctc ctgtgtctcc tctttcaccc agagagcttc tagcttttcc gccacttttt 120
 cggcatgata atttttgcct gatcctttct tttctctctc ttgatctctt ttcttgcatt 180
 cttcaaactt tgttttgaat ttctgtgcat tctcagcatt caggaagcgg atgg 234

<210> 266
 <211> 335
 <212> DNA
 <213> Homo sapien

<400> 266
 gtctcatca tcccagtttg aggcagtgtc ggagtgggga aggccttctt agaccataga 60
 ggttggaaga cgtgagaga tcatccagcc cagccccttg atgttacaga gcagaagaca 120
 gatgccaaa caggagaagg cacttgcca cggtcatacg gcaggttgcc acaaaaccaa 180
 gatggcagcc ctctctcagc gtgcctcact gccactccca gagccaggga gcccataaa 240
 accacatca tgtcttaaga gtatatctgg ctcttgacc agcaatcggc cctgggagcc 300
 accaggtggg aaaagcgcct ctgccagagt ccagg 335

<210> 267
 <211> 619
 <212> DNA
 <213> Homo sapien

<220>
 <221> misc_feature
 <222> (1)...(619)
 <223> n = A,T,C or G

<400> 267
 tggagctctg acgaagggat cggggaggtg ctggagaagg aagactgcat gcaggccctg 60

agcggccana	ttttcatggg	catggngtcc	tcccagttacc	aggccccggct	ggacatcgng	120
cgcttcattg	atgggcttgt	caacgcctgc	atccgctttg	tctacttctc	tttggaggat	180
gagctcaaaa	gcaaggtgtt	tgcanaaaaa	atgggcctgg	agacaggctg	gaactgccac	240
atctccctca	cacccaatgg	tgacatgcct	ggctccgaga	tccccccctc	cagccccagc	300
cacgcaggct	ccctgcatga	tgacctgaat	caggtgtccc	gagatgatgc	anaagggctc	360
ctcctcatgg	aggaggaggg	ccactcggac	ctcatcagct	tccagcctac	ggacagcgac	420
atccccagct	tcctggagga	ctccaaccgg	gccaagctgc	cccgggggat	ccaccaagtg	480
cgggccccacc	tgcagaacat	tgacaacgtg	cccctgctag	tgcccccttt	caccgactgc	540
accccanaga	ccatgtgtga	gatgataaag	atcatgcaan	agtacgggga	ggtgacctgc	600
tgcttgggca	nctctgcc					619

<210> 268

<211> 147

<212> DNA

<213> Homo sapien

<400> 268

cctataaccc	agacaccagc	atggacaaaa	ctcagttata	ctgaattcag	agacaaaatt	60
cagtgcact	cttctaccac	ttatttaggg	ttctacagca	tttctactgag	cagacttagt	120
tttttgtttt	tgttttacaa	acctttt				147

<210> 269

<211> 325

<212> DNA

<213> Homo sapien

<400> 269

ctgagctgta	ggaatgggtt	cttgggtacac	aagatagtat	tggtgagcta	gttttcgagc	60
tctgtgcaca	agcactctgt	aatcggggcc	catgccactg	tacaccaaac	ctatatgctt	120
ggtaattggg	tctactttgt	gtacacttcg	ctcatcatac	agaatggatt	tctgtttttt	180
ctcagttgct	aataccacac	catttgcagc	tttaattccc	acggacgggg	ctcctccagc	240
tacagcagcc	aaagcatatt	caatctggac	aagtttacca	gacgggctga	atgtagtcag	300
cgaaaagctg	tacccgcgct	ccgcc				325

<210> 270

<211> 428

<212> DNA

<213> Homo sapien

<400> 270

aaacatatgg	taaattaccg	agtgacacct	ctgggctaga	gacctctttt	gaggggagtt	60
tgcaaaactac	ggattcaatt	tctttaacag	ttatgaagtt	ctttaaagaa	cctgttttgt	120
attggggggg	tgtggtcacc	tgtgcttttc	tgagattttg	cccctacatc	taagttgttg	180
aatgcatgtg	tgtagagttg	tttatgggtc	ttccctttct	tcttagaagg	gtctatagta	240
atatccccctg	ccttatccct	agtagtacta	atttgtgttt	tcttacttct	tgacaggcaa	300
acacatcaga	gcataagtgg	ttcctaattg	caagctgacc	tcccttgatc	tctgtcttct	360
acaggatatt	gacatgggac	ttctttatta	ccttttcagt	tactgatac	cttcaaatag	420
ctttattt						428

<210> 271

<211> 206

<212> DNA

<213> Homo sapien

<220>
 <221> misc_feature
 <222> (1)...(206)
 <223> n = A,T,C or G

<400> 271
 cgtcccggag ccacacgngg ncatggctgg canagcgctc tgcattgctgg ggctggctcct 60
 ggcttctgctg tctccagct ctgctgagga gtacgtgggc ctgtctgcaa accagtgngc 120
 cgtgccagcc aaggacaggg tggactgagg ctacccccat gtcaccccca aggagtgcac 180
 caaccggggc tctgctttg actcca 206

<210> 272
 <211> 83
 <212> DNA
 <213> Homo sapien

<400> 272
 ctggttccc tgagaactca acaatgcctt ttctgaggg ccttctcga tcatccaaa 60
 tgactacagc cctctctacc tgg 83

<210> 273
 <211> 472
 <212> DNA
 <213> Homo sapien

<400> 273
 ctggagaagg tgtgcagggg aaaccctgct gatgtcaccg aggccagggt gtctttctac 60
 tcgggacact cttccttttg gatgtactgc atgggtgttct tggcgctgta tgtgcaggca 120
 cgactctgtt ggaagtgggc acggtctgctg cgaaccacag tccagttctt cctgggtggc 180
 ttgcccctct acgtgggcta cccccgctg tctgattaca aacaccactg gagcgatgtc 240
 cttgttggcc tctgacaggg ggcactgggtg gctgccctca ctgtctgcta catctcagac 300
 ttcttcaaag cccgaccccc acagcactgt ctgaaggagg aggagctgga acggaagccc 360
 agcctgtcac tgacgttgac cctgggcgag gctgaccaca accactatgg ataccgcac 420
 tctctctctt gaggcggac cccgcccagg cagggagctg ctgtgagtc ag 472

<210> 274
 <211> 205
 <212> DNA
 <213> Homo sapien

<400> 274
 ccaggcggcc cgaggactta cggtcggcac ttctctgttc tcccggtgca gcggtgtggtg 60
 tcgctgcat gggctgtacc tggatggtgt gtccaccatc gacacggagg ggctggattt 120
 gtttctcagg caatcctgta ttttaatttt agatgtattt cctgaagcat atttttcata 180
 gaatgtagcg tgtaaatagc ttttt 205

<210> 275
 <211> 308
 <212> DNA
 <213> Homo sapien

<400> 275
 ctctctgccc tccccaccga catcatgctc cagttccagc ttggatttac actgggcaac 60
 gtgggttgaa tgtatctggc tcagaactat gatatacaca acctgggctaa aaaacttgaa 120

aaacatacaa	taattttttat	tatggaaatt	aatctttaca	tacaaaatca	gctacgtaat	60
tttacttaca	aaacaataaa	aactgttctt	tactgtggca	acaaaagaag	cattttgaca	120
aatgaaaaaa	attaatgcaa	acaaattaaa	acaatgcttt	tctttttact	tgtttcactg	180
tctcttctat	ttattttcta	tgatcatttg	acacaaacat	ggattacttt	gatatctact	240
gaaacataaa	tgataagggt	cttaaagggt	gaattaaaag	tctgggtggt	caatatttta	300
gaagctgaat	aaacaaaacg	aaattggggg	ttgtgattac	agaggattta	tcattttttc	360
cctttgtcca	tatgaaaata	tataatagaa	aattaccac	gggaaaacat	tttt	414

<211> 262

<213> Hom

<213> Homo sapien

ccaccatgcc	tggcctgctt	caatTTTTTt	atgccacttt	gtaaacggca	cttaattatg	60
gaaaatagga	aaaagcaaaa	ctaaaataag	gaagaggata	tatatataac	ttttcacaat	120
ctctttttctg	atccccctta	gatgccagtt	caaccaggac	cacacacaga	tttcatttta	180
tttgtagagt	atatgaaaag	atttaatatg	ctcatgcatt	ttatttttacg	tatactgatt	240
tctacgtttt	gactgactat	tt				262

<211> 349

<212> DNA

<213> Home

<213> Homo sapien

ctgtgaccgc	ggtgcatcag	tggatatagt	tgtgtctccc	catggggggt	taacagtctc	60
tgcccaagac	cgttttctga	taatggctgc	agaaatggaa	cagtcattctg	gcacaggccc	120
agcagaatta	actcagtttt	ggaaagaagt	tcccagaaac	aaagtgatgg	aacatagggt	180
aagatgccat	actgttgaaa	gcagtaaac	aaacactctt	acgttaaaaag	acaatgcttt	240
caatatgtca	gataaaacca	gtgaagatat	atgtctacaa	ctcagtcggt	tactagaaag	300
caataggaag	cttgaagacc	aagttcagcg	ttgtattctgg	ttccagcag		349

<211> 381

<212> DNA

<213> Hom

<221> misc feature

$\langle 222 \rangle$ (1) ... (381)

<223> n = A, T, C o

aaacactaaa	tgaagcttct	cacaatttct	aattataaac	aaaaggtga	aaacagtatg	60
ggaaacaaag	tttcaaaaca	aagaaaagtt	gagtaaaagg	tgccccctct	atggtctatc	120
tgaaagaaac	attttactca	gagaggcaaa	cattttctgat	ctaggagtaa	gtttccact	180
cactttgcaa	ggaccctactc	attctgcana	aagacctaca	agtctttctg	gtctcaattg	240
caaagtacgt	gaaaatgtgt	atgaaagatc	taaaagctaa	atattagaat	aaggctaatt	300
gaaatcaaaa	ttgtgtgctg	gtctaaatat	acatcttcgg	cttcttctct	tttagtaagt	360
atttttattt	caqatqtatt	t				381

<210> 283

<211> 543
 <212> DNA
 <213> Homo sapien

<400> 283
 aatatagctc ctccctaccc ccaacaatgg accctgcccc ttgcctccca gttccttgat 60
 cttcctagggt tccacaactc tcttttttcc ttttagtttta ttccctccag ccaaacctct 120
 cttattcaat attttgagcc aatgggggag ttatgtagat ttttttccct acacattagc 180
 tggccctttt tatgaccaat gactcataag gcaagatgtg tgggtggcatc ttcggacagg 240
 cagcaggctt taatagggca gcctggggtt gtggaggcaa gcaaagctaa ttggcatgcg 300
 tgggaatcaa accccaggcc ctgggctcat tagcccatgg tcaaaacaac tgagccagag 360
 gaggtataaa tttgcccagg aatatcagta gttcctttat tagaagaaaa tggctgatat 420
 ggaagttggg gaatctgaat tgccagagaa tcttggggaag agtaataagc tcttagtctc 480
 aacaaaaaagt gttttttcat ctcagcgcgt aaaggggtgct atatgggaac aaagaagtat 540
 ttt 543

<210> 284
 <211> 147
 <212> DNA
 <213> Homo sapien

<400> 284
 aaactggtat tttatctttg attctccttc agccctcacc cctggttctc atctttcttg 60
 atcaacatct tttcttgctt ctgtcccctt ctctcatctc ttagctcccc tccaacctgg 120
 ggggcagtgg tgtggagaag ccacagg 147

<210> 285
 <211> 316
 <212> DNA
 <213> Homo sapien

<400> 285
 cgcccgaggt ctggcttcac tctactccc tctctgctcg cagcacgtcg gccgccagct 60
 ctttgatgtg ttcccaggcc cgctgcacat gggcagattc caccgtgcga gaacagatgg 120
 caaagcgcag gacaaaactg tccctgaggt gacatggaac caagtggatt tttttggcac 180
 tgtttattct ttgcagaaga gcttcattca ctttgttggg acccttttagc cgaaagcaga 240
 caagccccag aatgacttcc acacagattt caaagcgggg atcctggcgc accagtgact 300
 caaactcatg ggacag 316

<210> 286
 <211> 322
 <212> DNA
 <213> Homo sapien

<400> 286
 cctggggagc ccttttagtg ggtgggacct caggcagacc cccaaaccaa agggagccag 60
 atgcccaggt tcaagtcatt agtgatatgt ggcagggctg acagagaaat aatcctggag 120
 gtctccaaag ctgctgggaa tggaaatggc atgaaaagcg caggagtggg cagggtgtgg 180
 tgggtgatgg tggcctcact cagagtggac caaggcccca gtccttgcc caaaaccaa 240
 gcccttgggc ccgaagtttt tagcataaca tcctttgcag taaatctcgc catccttgtc 300
 tgccaggggtg gttgactcaa gg 322

<210> 287
 <211> 364

<212> DNA
<213> Homo sapien

<400> 287
ctgcccacgc tcaaaccaat tctggctgat atcgagtacc tgcaggacca gcacctcctg 60
ctcacagtca agtccatgga tggctatgaa tcctatgggg agtgtgtggt tgcaactcaa 120
tccatgatcg gcagcacggc ccaacagttc ctgaccttcc tatcccaccg tggcgaggag 180
acaggcaata tcagaggctc catgaagggtg cgggtgcccc cggagcgcct gggcaccctg 240
gagcggctct acgagtggat cagcattgat aaggatgagg caggagcaaa gagcaaagcc 300
ccctctgtgt cccgagggag ccaggagccc aggtcaggga gccgcaagcc agccttcaca 360
gagg 364

<210> 288
<211> 261
<212> DNA
<213> Homo sapien

<400> 288
aaaattataa ctactcattc tttcttttagc cttagttaat ttgagcagaa gccacaacaa 60
gcaaaccaca ataaatttag aattggcaga aatccacatt aactcctctt cccaagtctt 120
cacactacta ccattttacag ttgtagggtt gtaatgtata attatgtaat gcagaaacta 180
gctttgactt gtgtaacgat gcactgtcaa agtaagcaaa gtaagaattg aaattccaca 240
ttcccagaat ttaacactca g 261

<210> 289
<211> 261
<212> DNA
<213> Homo sapien

<400> 289
ctgagtgtta aattctggga atgtggaatt tcaattotta ctttgottac tttgacagtg 60
catcgttaca caagtcaaag ctagttttctg cattacataa ttatacatta caaacctaca 120
actgtaaatg gtagtagtgt ggaaacttgg gaagaggagt taatgtggat ttctgccaat 180
tctaaattta ttgtggtttg cttgttgtgg cttctgctca aattaactaa ggctaaagaa 240
agaatgagta gttataattt t 261

<210> 290
<211> 92
<212> DNA
<213> Homo sapien

<400> 290
ccactacccg aacttacagg tgccaaaaga agaaagggta taaacggaga ccacctatca 60
ctcatcagaa ctaggatca tcacattcct tt 92

<210> 291
<211> 287
<212> DNA
<213> Homo sapien

<400> 291
ccatggctcc gctcagggcc ccggtcacct ccgagtcact ctgttccttg actgtctttg 60
tgtttctgta cctcaaggca ctgaagctgg aggactctgt ccatgcctgt gtcaccctcg 120
tgtgggagcc tctgggctcg gcaggtccac atttcatgag ctgaggcgtg ggccagggcc 180

```
<210> 292
<211> 270
<212> DNA
<213> Homo sapien
```

```
<210> 293
<211> 333
<212> DNA
<213> Homo sapien
```

```
<210> 294
<211> 123
<212> DNA
<213> Homo sapien
```

```
<210> 295
<211> 311
<212> DNA
<213> Homo sapien
```

<210>	296
<211>	241
<212>	DNA

<400> 296

<210> 297

<211> 295

<212> DNA

<213> Homo sapien

<400> 297

aaacacaaga	tgaaaaataact	ctgttctgtc	caaagcatca	cctaattggtg	tgaggcatct	60
cacttagctg	tggagaagtc	cttggaatta	gatctcagaa	agacagcttt	aagacagtaa	120
aaccttttgg	caatgggcta	attgccttaa	aagaagagtt	ctacctgaaa	gaccttgcag	180
gtggagaaat	tgtcctacaa	agattcttgg	atatgttagt	ggagataact	gacatgggta	240
gctgtgggtc	aaccaggaac	tgtcaacaac	ctgatctctg	caaaaccagg	atgga	295

<210> 298

<211> 347

<212> DNA

<213> Homo sapien

<400> 298

ccttcaggca	agaggcaaag	atccagtgga	atatgggaga	atggtggagg	60
accaacacct	gctaccccag	agagcttttc	taaaaaaagc	aagaaagcag	120
tattcaccct	gcagaagaca	cggaaaggtac	tgagtttgag	ccagaggggac	180
tgtaaaagaaa	gggtttgctg	acatcccagc	aggaaagact	agcccatata	240
aacaaccatg	gcaactcgga	ccagcccccg	cctggctgca	cagaagttag	300
actgagtctc	ggcaaagaaa	atcttgcaga	gtcctccaaa	ccaacag	347

<210> 299

<211> 268

<212> DNA

<213> Homo sapien

<400> 299

aaaaagtaaa	catgaaaaca	tcacgaattg	taccatgatt	caagaataac	ttttgtaata	60
gaaaacacat	gaccttttgc	agtatagtgt	gataccgaag	taaaagtgaa	agaaataaat	120
gcaggaaagt	ttaagtggat	gtaagttttt	ataaggaaag	taataagagg	aggctgcttt	180
tgaaggtcct	ttgatcttcc	atgatgataa	tatcgttgca	aagttcttta	acttgatttc	240
aagtaattag	cagttgacca	cttggttt				268

 $\langle 210 \rangle$ 300

<211> 185

<212> DNA

<213> Homo sapien

<400> 300

aaattggaga	aggaagtttt	cctgaagagc	cagaatcctt	gctaagtcac	ttagatccaa	60
ctgaccactc	ttattttctgt	caaaaatcct	catcatgggtg	ccgggtgtatt	cttccagttt	120

agcctcagaa atggcctttc tgtggtgaag aaagaggctc cggaggaagt tgcggagctc 180
agcag 185

<210> 301
<211> 75
<212> DNA
<213> Homo sapien

<400> 301
aaaattggaa agtgggataa gaaatctaaa gtaaccagct tatctttgaa acaatattat 60
tttgaaattg gcttt 75

<210> 302
<211> 247
<212> DNA
<213> Homo sapien

<220>
<221> misc_feature
<222> (1)...(247)
<223> n = A,T,C or G

<400> 302
ccatgtttctc tgaattgggt gcagaagaca agggcagagt ggctgcggcc cctattacct 60
ttgtagcagc cacatcagaa agcagaagaa aacagtatct ctgaaggcat tgtttgaggt 120
tgatctcagc actgaacgat ttcaagccct acgcaccana acagaaggag ggtggaggaa 180
gtgatcanag ggaacgagct gtaggtttgc anaaatgtgt gaaaccaaaa tgatcactgc 240
ctacttg 247

<210> 303
<211> 535
<212> DNA
<213> Homo sapien

<400> 303
ctgcttcaga ggaaatcact gaaaaataaa gaaaaacat ccatgcatgg ctgcatccag 60
tgtacctgta atcctgaaga aaaggctccta attccttcca tgctgaaatg ctagcttttg 120
tttcagagag agactttatt gcaactgtga ccaccgtcac tggtagcac tgctgttcgg 180
ccccagcgg acttaaaaga ctggaatgtg gtagtgccgg tcgttctcgg tcagcaggga 240
gatctccggc cagtccttga gaggtcctc tgggtagcag acttcaaagt ctctggagtt 300
aaacttgaac agtctgaaca cttttatctt tacttcaagg gagtatccaa gtataaacat 360
atcaatctgc tctagtccac atgtgtcggc tacagaattc aggtgattca tcatgaagct 420
caaaggatca gaggatgtct ccctggaaaa caggagtcta aaaagactgg gaatgacctt 480
tttagtcttc atttgttcat aaacttcagt gacttgatac agcatgatga acttt 535

<210> 304
<211> 522
<212> DNA
<213> Homo sapien

<400> 304
ccgcgctcgg tctacaatca cgttttatta ttggctcgtc tagtcatggg atagagaagg 60
taaatagcaa aatagaaaga aaagggggaa aaggtagaag gcaaggggaa aactattgg 120
tttagatctt tatcctgggtc ctgtcaatga tcaggtaatt ggaaggatca aaattaggcc 180

```
<210> 305
<211> 165
<212> DNA
<213> Homo sapien
```

```
<210> 306
<211> 294
<212> DNA
<213> Homo sapien
```

```
<210> 307
<211> 181
<212> DNA
<213> Homo sapien
```

```
<210> 308
<211> 179
<212> DNA
<213> Homo sapien
```

<400> 308						
aaggtctgagg	actgctggga	gctcagatca	gcccggagct	actggctcat	gggcagccaa	60
aaaatactgg	atctgctgaa	cgaaggctca	gcccgagatc	tccgcagtct	tcagcgcatt	120
ggccccgaaga	aggcccanct	aatcgtgggc	tggcgggagc	tccacggccc	cttcagcca	179

<210> 309
 <211> 129
 <212> DNA
 <213> Homo sapien

<220>
 <221> misc_feature
 <222> (1)...(129)
 <223> n = A,T,C or G

<400> 309	
ctgcccgttt gcccgtagct gactcagntt cctcatcttc atctccatcc tcttctcac	60
catcaccttc ttcttctctc tctcttctc cccaccttc ttctcttctc tcgtctacct	120
cattgtcag	129

<210> 310
 <211> 390
 <212> DNA
 <213> Homo sapien

<400> 310	
tgaggctggg ggagagccgt ggtccctgag gatgggtcag agctaaactc cttcctggcc	60
tgagagtcag ctctctgccc tgtgtacttc cggggccagg gctgccccta atctctgtag	120
gaaccgtggg atgtctgcat gttgcccctt tctcttttcc cctttcctgt cccaccatac	180
gagcacctcc agcctgaaca gaagctctta ctctttccta ttccagtgtt acctgtgtgc	240
ttgggtctgtt tgactttacg cccatctcag gacacttccg tagactgttt aggttcccct	300
gtcaaataac agttaccac tcgggtcccag ttttgttgcc ccagaaaggg atgttattat	360
ccttgggggc tcccagggca aggggttaagg	390

<210> 311
 <211> 355
 <212> DNA
 <213> Homo sapien

<220>
 <221> misc_feature
 <222> (1)...(355)
 <223> n = A,T,C or G

<400> 311	
cctctctgtg ctgctgaagg cagatcgctt gttccacacc agctaccact cccaggcagt	60
gcataatccg ctgttgagaa atgccgtgtc tagattgtgg acaagagcct gogtgattat	120
gctatangga naaaaattct tcgagtcca ccnancctc tctaaacatt tggctcactc	180
aaaacaaaaa gncaccaatc ttantactgc tgaacttcat ttatgttnacc taacattaac	240
cntcgtagga aaaccaaata gccctctcgt ncangatatg ttgctaaagg actacctgt	300
tcaacacaac ggctccggtg tgtgaactcc tgtttgggtg attcccctac tctca	355

<210> 312
 <211> 498
 <212> DNA
 <213> Homo sapien

<400> 312

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ccattctttt gaatctaata tattatcaat agcatcctcc ataatatctt tgataaaaagg      60
tgtccaccga gagagctgaa aagtttcttc tgcagaccga tcctttctta acggtttgcc      120
ttgttgagat tggggaacaa tgggaacacc aaggtaactc cagttacgaa tcatgtcact      180
ctcattttct atctttacat tctggatcaa cctgtccaaa ttttcttcgg tagttccatt      240
aatactgaag atataaagta gaattgctct tattttatca caattatcat gatttttgtt      300
gagtagaact ggaaggagta ctgcacatgga atctttcacc ttctgtcctt ctgcatcagt      360
tccaagtgcc aggtcctgtt cagttttgca gagcttttct atattaagct tgaacttatt      420
catgcaatct tctgctaagt taagatggac aacttgctta gtaatctgtt ttcggaaata      480
gggcatcttt ttcacatcag

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<210> 313
<211> 653
<212> DNA
<213> Homo sapien

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<400> 313
aaacttatca gattttttta agttaggtaa tttcaatcca cagtggctcc atatggttaa      60
aaaaacaaaa acaaaaacgc atttaaggat acacgaagca gtgaaaacaa agccccagta      120
ttttcgctaa agtactggaa atacctgttt ctaaaaacag ctttatattt gtccactgcc      180
tagaatagct ctcacccaaa cctcaaaaat aagagcagat agatttttaga agcaagaaaa      240
ggtaaacagt gcccatatta tttgagactg gctctgctgc cctccctaag ccagtttaca      300
ttctttgaga ttcttgaggt ggggtgagtc gggctgaaga ctgcacaggc catgtcccct      360
gctccaacta ttctcagaa cgtcccagggt ggagggagtg gctgtcgat tttcactcat      420
tccatggagc tctgtgtaca tgaaaattcc tccaagtgtg gcttttgtcg aattcagaga      480
tacagcaagc cacgcataaa acatggagtg tagagcactg gtgtacctag cttagaaaca      540
ccctcgggtga atgtggtact gtggctcgaa aggaagcaag ggacaggacc caggagactg      600
ggcgccagg ctctcggagt tccacacaca cctgtgaagc ccggccagca cag

```

```

<210> 314
<211> 513
<212> DNA
<213> Homo sapien

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```

<400> 314
ctggaagatt ttgctgcatt tggcattata ctgtaattta cagtatacaa catctgggga      60
ctcagtacta tcttagcaca gactaacttc tccactccg tcagaggtgg cagggtggcgg      120
gtcgggtggg agggcctttt ctcccataa atgcctgaac tttaatttat accatataag      180
aatcagtgta aaggtaaaca acaaggttta tgtaactcta ttataaattt tgcatttttt      240
ttctctgtga catatacaag tatatttttg tttttggagc tataaattat ttaatttagc      300
aatcttcaaa gctcataaat ttcaactttt caaataagaa attttaactt caaataagaa      360
gtctaggact ttatggctat taattttact atcaaaatat ccaagggact ccattcaatg      420
taatagttat aattcttcta aatatcattt gaataattct ttgtggacgc tagactcaag      480
actatgctac atccaaacag tacatctata acc

```

```

<210> 315
<211> 222
<212> DNA
<213> Homo sapien

```

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<220>
<221> misc_feature
<222> (1)...(222)
<223> n = A,T,C or G

```

<400> 315

atttatattc	aaggngatctc	aaagaaagca	ttttcatttc	actgcacatc	tagagaaaaa	60
caaaaataga	aaatcttcta	gtccatccta	atctgaatgg	tgctgtttct	atattgggtca	120
ttgccttgca	aacaggagct	ccacaaaagc	caggaagaga	gactgcctcc	ttggctgaaa	180
gagtcctttc	aggaaggtgg	actgcattgg	tttgatatgt	tt		222

<210> 316

<211> 1633

<212> DNA

<213> Homo sapiens

<400> 316

cgtggaggca	gctagcgcga	ggctggggag	cgctgagccg	cgcgctcgtgc	cctgcgctgc	60
ccagactagc	gaacaataca	gtcgggatgg	ctaaagggtga	ccccaagaaa	ccaaagggca	120
agacgtccgc	ttatgccttc	tttgtgcaga	catgcagaga	agaacataag	aagaaaaacc	180
cagaggtccc	tgtcaatttt	gcggaatttt	ccaagaagtg	ctctgagagg	tggaagacgg	240
tgtccgggaa	agagaaatcc	aaatttgatg	aaatggcaaa	ggcagataaa	gtgcgctatg	300
atcgggaaat	gaaggattat	ggaccagcta	agggaggcaa	gaagaagaag	gatcctaata	360
ctcccaaaag	gccaccgtct	ggattcttcc	tggtctgttc	agaattccgc	cccaagatca	420
aatccacaaa	ccccggcatc	tctattggag	acgtggcaaa	aaagctgggt	gagatgtgga	480
ataatttaaa	tgacagtga	aagcagcctt	acatcactaa	ggcggcaaa	ctgaaggaga	540
agtatgagaa	ggatgttgct	gactataagt	cgaaaggaaa	gtttgatggg	gcaaaggggc	600
ctgctaaaagt	tgcccgga	aaggtggaag	aggaagatga	agaacaggag	gaggaagaag	660
aggaggagga	ggaggaggag	gatgaataaa	gaaactgttt	atctgtctcc	ttgtgaatac	720
ttagagtagg	ggagcgccgt	aattgacaca	tctcttattt	gagaagtgtc	tggtgccctc	780
attaggttta	attacaaaat	ttgatcacga	tcatattgta	gtctctcaaa	gtgctctaga	840
aattgtcagt	ggtttacatg	aagtggccat	gggtgtctgg	agcacctga	aactgtatca	900
aagtgtgaca	tatttccaaa	catttttaaa	atgaaaaggc	actctcgtgt	tctcctcact	960
ctgtgcactt	tgctgttggt	gtgacaaggc	atttaaagat	gtttctggca	ttttcttttt	1020
atgtgtaagg	tggtggtaac	tatggttatt	ggctagaaat	cctgagtttt	caactgtata	1080
tatctatagt	ttgtaaaaag	aacaaaacaa	ccgagacaaa	cccttgatgc	tccttgctcg	1140
gcgttgaggc	tgtggggaag	atgccttttg	ggagaggctg	tagctcaggg	cgtgcactgt	1200
gaggctggac	ctgttgactc	tgcaaggggc	atccatttag	cttcagggtg	tcttgtttct	1260
gtatatagt	acatagcatt	ctgctgccat	cttagctgtg	gacaaagggg	ggtcagctgg	1320
catgagaata	ttttttttta	agtgcggtag	tttttaaact	gtttgttttt	aaacaaacta	1380
tagaactctt	cattgtcagc	aaagcaaaga	gtcactgcac	caatgaaagt	tcaagaacct	1440
cctgtactta	aacacgattc	gcaacgtttc	gttatttttt	ttgtatgttt	agaatgctga	1500
aatgtttttg	aagttaaata	aacagtatta	cattttttaga	actcttctct	actataacag	1560
tcaattttctg	actcacagca	gtgaacaaac	ccccactccg	ttgtatttgg	agactggcct	1620
ccctataaat	gtg					1633

<210> 317

<211> 4235

<212> DNA

<213> Homo sapiens

<400> 317

gaatccaagg	gggccagttc	ctgccgtctg	ctcttctgcc	tcttgatctc	cgccaccgtc	60
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ttttctaaag aaagctgtgt tcttctgttg acccagacga atagggcaca gccctgtaac 1920
tgacagtgcc ttctgtcatt gggaatgaaa taaattatta cgagaaaggg acttgctcta 1980
actggtttga ggccttacag ttttgtatct acatttttcc cctcctgggg tttgcgggga 2040
cagggacaga actacaggag tcatgggaaa gaaaattctg gcttcactac tgctcactgc 2100
tcactttctg atcactctga tacttttttt tttttttttt ttttgcaacc tgataccttg 2160
aaaagcttct atgtgtctct ccttttgttg cctggcagct gtctaggatg atcactgatt 2220

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actatcttact aagtagccac atgcaaataa aagttgtttg gtaaaatgga aaaaaaaaaa 2280

<210> 322

<211> 1398

<212> DNA

<213> Homo sapiens

<400> 322

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ctaaccctaa	ggaattgaaa	ggaaccactc	attcacttct	agacgacaaa	atgcaaaaaa	180
ggaggccaaa	gacttttgga	atggatatga	aagcatacct	gagatctatg	atcccacatc	240
tgaatctgg	aatgaaatct	tccaagtcca	aggatgtact	ttctgctgct	gaagtaatgc	300
aatggtctca	atctctggaa	aaacttcttg	ccaacccaaac	tgggtcaaat	gtctttggaa	360
gtttcctaaa	gtctgaattc	agtgaggaga	atattgagtt	ctggctggct	tgtgaagact	420
ataagaaaac	agagtctgat	cttttgccct	gtaaagcaga	agagatatat	aaagcatttg	480
tgcattcaga	tgtgtctaaa	caaatacaata	ttgacttccg	cactcgagaa	tctacagcca	540
agaagattaa	agcaccaacc	cccacgtgtt	ttgatgaagc	acaaaaagtc	atatatactc	600
ttatggaaaa	ggactcttat	cccagggtcc	tcaaatcaga	tatttactta	aatcttctaa	660
atgacctgca	ggctaatagc	ctaaagtgc	tgggtccctgg	ctgaaggga	ttaacagata	720
gtatcaaggc	acgaaggaa	gtgccagtat	ggctccctgg	gtgaacagct	tggccttttt	780
tgggtgtctt	gacaggccaa	gaagaacaaa	tgactcagaa	tggattaaca	tgaaagttat	840
ccaggcgag	agttgaagaa	gcataagcaa	gacaaaaaca	gagagaccgc	agaaggagga	900
agatactgtg	gtactgtcat	aaaaaacagt	ggagctctgt	attagaaagc	ccctcagaac	960
tgggaaggcc	aggtaactct	agttacacag	aaactgtgac	taaagtctat	gaaactgatt	1020
acaacaggct	gtaagaatca	aagtcaactg	acatctatgc	tacatatatt	tatatagttt	1080
gtactgagct	attgaagtcc	cattaactta	aagtatatgt	tttcaaattg	ccattgctac	1140
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tgggtgcta	aataaatgtg	gattttgtat	taaaatatat	agaagcaatt	tctgtttaca	1320
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<210> 323

<211> 1316

<212> DNA

<213> Homo sapiens

<400> 323

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gtgaaagaag	cagtgaagg	ggccattgat	gcaggatata	ggcacattga	ctgtgcctat	180
gtctatcaga	atgaacatga	agtgggggaa	gcatccaag	agaagatcca	agagaaggct	240
gtgaagcggg	aggacctgtt	catcgtcagc	aagttgtggc	ccactttctt	tgagagaccc	300
cttgtgagga	aagcctttga	gaagaccctc	aaggacctga	agctgagcta	tctggacgtc	360
tattcttatt	actggccaca	gggattcaag	tctggggatg	accttttccc	caaagatgat	420
aaaggtaatg	ccatcggtgg	aaaagcaacg	ttcttggatg	cctggggaggc	catggaggag	480
ctggtggatg	aggggctgg	gaaagccctt	ggggtctcca	atttcagcca	cttcagatc	540
gagaagctct	tgaacaaacc	tggactgaaa	tataaaccag	tgactaacca	ggttgagtgt	600
caccataacc	tcacacagga	gaaactgata	cagtactgcc	actccaagg	catcaccgtt	660
acggcctaca	gccccctggg	ctctccggat	agaccttggg	ccaagccaga	agacccttcc	720
ctgctggagg	atoccaaag	taaggagatt	gctgcaaagc	acaaaaaac	cgcagcccag	780
gttctgatcc	gtttccatat	ccagaggaat	gtgattgtca	tccccaagtc	tgtgacacca	840
gcacgcattg	ttgagaacat	tcagggtctt	gacttttaaa	tgagtgatga	ggagatggca	900

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accatactca gcttcaacag aaactggagg gcctgtaacg tgttgcaatc ctctcatttg 960
gaagactatc ccttcaatgc agaattattga gggtgaatct cctgggtgaga ttatacagga 1020
gattctcttt ctctgctgaa gtgtgactac ctccactcat gtcccatttt agccaagctt 1080
atttaagatc acagtgaact tagtcctgtt atagacgaga atcgaggtgc tgttttagac 1140
atttatttct gtatgttcaa ctaggatcag aatatcacag aaaagcatgg cttgaataag 1200
gaaatgacaa ttttttccac ttatctgacg agaacaaatg tttattaagc atcagaaact 1260
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<210> 324

<211> 200

<212> PRT

<213> Homo sapiens

<400> 324

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Met Ala Lys Gly Asp Pro Lys Lys Pro Lys Gly Lys Thr Ser Ala Tyr
      5                      10                      15

Ala Phe Phe Val Gln Thr Cys Arg Glu Glu His Lys Lys Lys Asn Pro
      20                      25                      30

Glu Val Pro Val Asn Phe Ala Glu Phe Ser Lys Lys Cys Ser Glu Arg
      35                      40                      45

Trp Lys Thr Val Ser Gly Lys Glu Lys Ser Lys Phe Asp Glu Met Ala
      50                      55                      60

Lys Ala Asp Lys Val Arg Tyr Asp Arg Glu Met Lys Asp Tyr Gly Pro
      65                      70                      75                      80

Ala Lys Gly Gly Lys Lys Lys Lys Asp Pro Asn Ala Pro Lys Arg Pro
      85                      90                      95

Pro Ser Gly Phe Phe Leu Phe Cys Ser Glu Phe Arg Pro Lys Ile Lys
      100                     105                     110

Ser Thr Asn Pro Gly Ile Ser Ile Gly Asp Val Ala Lys Lys Leu Gly
      115                     120                     125

Glu Met Trp Asn Asn Leu Asn Asp Ser Glu Lys Gln Pro Tyr Ile Thr
      130                     135                     140

Lys Ala Ala Lys Leu Lys Glu Lys Tyr Glu Lys Asp Val Ala Asp Tyr
      145                     150                     155                     160

Lys Ser Lys Gly Lys Phe Asp Gly Ala Lys Gly Pro Ala Lys Val Ala
      165                     170                     175

Arg Lys Lys Val Glu Glu Glu Asp Glu Glu Gln Glu Glu Glu Glu Glu
      180                     185                     190

Glu Glu Glu Glu Glu Glu Asp Glu
      195                     200

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006280" 636F5960

<210> 325
 <211> 263
 <212> PRT
 <213> Homo sapiens

<400> 325
 Met Phe Arg Asn Gln Tyr Asp Asn Asp Val Thr Val Trp Ser Pro Gln
 5 10 15
 Gly Arg Ile His Gln Ile Glu Tyr Ala Met Glu Ala Val Lys Gln Gly
 20 25 30
 Ser Ala Thr Val Gly Leu Lys Ser Lys Thr His Ala Val Leu Val Ala
 35 40 45
 Leu Lys Arg Ala Gln Ser Glu Leu Ala Ala His Gln Lys Lys Ile Leu
 50 55 60
 His Val Asp Asn His Ile Gly Ile Ser Ile Ala Gly Leu Thr Ala Asp
 65 70 75 80
 Ala Arg Leu Leu Cys Asn Phe Met Arg Gln Glu Cys Leu Asp Ser Arg
 85 90 95
 Phe Val Phe Asp Arg Pro Leu Pro Val Ser Arg Leu Val Ser Leu Ile
 100 105 110
 Gly Ser Lys Thr Gln Ile Pro Thr Gln Arg Tyr Gly Arg Arg Pro Tyr
 115 120 125
 Gly Val Gly Leu Leu Ile Ala Gly Tyr Asp Asp Met Gly Pro His Ile
 130 135 140
 Phe Gln Thr Cys Pro Ser Ala Asn Tyr Phe Asp Cys Arg Ala Met Ser
 145 150 155 160
 Ile Gly Ala Arg Ser Gln Ser Ala Arg Thr Tyr Leu Glu Arg His Met
 165 170 175
 Ser Glu Phe Met Glu Cys Asn Leu Asn Glu Leu Val Lys His Gly Leu
 180 185 190
 Arg Ala Leu Arg Glu Thr Leu Pro Ala Glu Gln Asp Leu Thr Thr Lys
 195 200 205
 Asn Val Ser Ile Gly Ile Val Gly Lys Asp Leu Glu Phe Thr Ile Tyr
 210 215 220
 Asp Asp Asp Asp Val Ser Pro Phe Leu Glu Gly Leu Glu Glu Arg Pro
 225 230 235 240
 Gln Arg Lys Ala Gln Pro Ala Gln Pro Ala Asp Glu Pro Ala Glu Lys
 245 250 255

006280" E3E5960

Glu Gly Leu Val Leu Thr Gln Lys Val Ser Asn Ser Gly Ile Thr Arg
225 230 235 240

Val Glu Lys Ala Lys Ile Gly Leu Ile Gln Phe Cys Leu Ser Ala Pro
 245 250 255
 Lys Thr Asp Met Asp Asn Gln Ile Val Val Ser Asp Tyr Ala Gln Met
 260 265 270
 Asp Arg Val Leu Arg Glu Glu Arg Ala Tyr Ile Leu Asn Leu Val Lys
 275 280 285
 Gln Ile Lys Lys Thr Gly Cys Asn Val Leu Leu Ile Gln Lys Ser Ile
 290 295 300
 Leu Arg Asp Ala Leu Ser Asp Leu Ala Leu His Phe Leu Asn Lys Met
 305 310 315 320
 Lys Ile Met Val Ile Lys Asp Ile Glu Arg Glu Asp Ile Glu Phe Ile
 325 330 335
 Cys Lys Thr Ile Gly Thr Lys Pro Val Ala His Ile Asp Gln Phe Thr
 340 345 350
 Ala Asp Met Leu Gly Ser Ala Glu Leu Ala Glu Glu Val Asn Leu Asn
 355 360 365
 Gly Ser Gly Lys Leu Leu Lys Ile Thr Gly Cys Ala Ser Pro Gly Lys
 370 375 380
 Thr Val Thr Ile Val Val Arg Gly Ser Asn Lys Leu Val Ile Glu Glu
 385 390 395 400
 Ala Glu Arg Ser Ile His Asp Ala Leu Cys Val Ile Arg Cys Leu Val
 405 410 415
 Lys Lys Arg Ala Leu Ile Ala Gly Gly Gly Ala Pro Glu Ile Glu Leu
 420 425 430
 Ala Leu Arg Leu Thr Glu Tyr Ser Arg Thr Leu Ser Gly Met Glu Ser
 435 440 445
 Tyr Cys Val Arg Ala Phe Ala Asp Ala Met Glu Val Ile Pro Ser Thr
 450 455 460
 Leu Ala Glu Asn Ala Gly Leu Asn Pro Ile Ser Thr Val Thr Glu Leu
 465 470 475 480
 Arg Asn Arg His Ala Gln Gly Glu Lys Thr Ala Gly Ile Asn Val Arg
 485 490 495
 Lys Gly Gly Ile Ser Asn Ile Leu Glu Glu Leu Val Val Gln Pro Leu
 500 505 510
 Leu Val Ser Val Ser Ala Leu Thr Leu Ala Thr Glu Thr Val Arg Ser
 515 520 525

006230"EGF5960

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<210> 327
<211> 144
<212> PRT
<213> Homo sapiens
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Thr Ala Ala Leu Ile Phe Phe Ala Ile Trp His Ile Ile Ala Phe Asp
20 25 30

Glu Leu Lys Thr Asp Tyr Lys Asn Pro Ile Asp Gln Cys Asn Thr Leu
35 40 45

Asn Pro Leu Val Leu Pro Glu Tyr Leu Ile His Ala Phe Phe Cys Val
50 55 60

Met Phe Leu Cys Ala Ala Glu Trp Leu Thr Leu Gly Leu Asn Met Pro
65 70 75 80

Leu Leu Ala Tyr His Ile Trp Arg Tyr Met Ser Arg Pro Val Met Ser
85 90 95

Gly Pro Gly Leu Tyr Asp Pro Thr Thr Ile Met Asn Ala Asp Ile Leu
100 105 110

Ala Tyr Cys Gln Lys Glu Gly Trp Cys Lys Leu Ala Phe Tyr Leu Leu
115 120 125

Ala Phe Phe Tyr Tyr Leu Tyr Gly Met Ile Tyr Val Leu Val Ser Ser
130 135 140

```
<210> 328
<211> 138
<212> PRT
<213> Homo sapiens
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<400> 328
Met Pro Asn Phe Ser Gly Asn Trp Lys Ile Ile Arg Ser Glu Asn Phe
 5 10 15

Glu Glu Leu Leu Lys Val Leu Gly Val Asn Val Met Leu Arg Lys Ile
20 25 30

Ala Val Ala Ala Ala Ser Lys Pro Ala Val Glu Ile Lys Gln Glu Gly
35 40 45

Asp Leu Pro Glu Asp Val Lys Trp Ile Asp Ile Thr Pro Asp Met Met
145 150 155 160

Asn Val Ser Ser His Leu Asp Lys Ala Ser Val Met Arg Leu Thr Ile

	50					55					60				
Ser	Tyr	Leu	Arg	Val	Arg	Lys	Leu	Leu	Asp	Ala	Gly	Asp	Leu	Asp	Ile
65					70					75					80
Glu	Asp	Asp	Met	Lys	Ala	Gln	Met	Asn	Cys	Phe	Tyr	Leu	Lys	Ala	Leu
				85					90					95	
Asp	Gly	Phe	Val	Met	Val	Leu	Thr	Asp	Asp	Gly	Asp	Met	Ile	Tyr	Ile
			100					105					110		
Ser	Asp	Asn	Val	Asn	Lys	Tyr	Met	Gly	Leu	Thr	Gln	Phe	Glu	Leu	Thr
		115					120					125			
Gly	His	Ser	Val	Phe	Asp	Phe	Thr	His	Pro	Cys	Asp	His	Glu	Glu	Met
	130					135					140				
Arg	Glu	Met	Leu	Thr	His	Arg	Asn	Gly	Leu	Val	Lys	Lys	Gly	Lys	Glu
145					150					155					160
Gln	Asn	Thr	Gln	Arg	Ser	Phe	Phe	Leu	Arg	Met	Lys	Cys	Thr	Leu	Thr
				165					170					175	
Ser	Arg	Gly	Arg	Thr	Met	Asn	Ile	Lys	Ser	Ala	Thr	Trp	Lys	Val	Leu
			180					185					190		
His	Cys	Thr	Gly	His	Ile	His	Val	Tyr	Asp	Thr	Asn	Ser	Asn	Gln	Pro
		195					200					205			
Gln	Cys	Gly	Tyr	Lys	Lys	Pro	Pro	Met	Thr	Cys	Leu	Val	Leu	Ile	Cys
	210					215					220				
Glu	Pro	Ile	Pro	His	Pro	Ser	Asn	Ile	Glu	Ile	Pro	Leu	Asp	Ser	Lys
225					230					235					240
Thr	Phe	Leu	Ser	Arg	His	Ser	Leu	Asp	Met	Lys	Phe	Ser	Tyr	Cys	Asp
				245					250					255	
Glu	Arg	Ile	Thr	Glu	Leu	Met	Gly	Tyr	Glu	Pro	Glu	Glu	Leu	Leu	Gly
			260					265					270		
Arg	Ser	Ile	Tyr	Glu	Tyr	Tyr	His	Ala	Leu	Asp	Ser	Asp	His	Leu	Thr
		275					280					285			
Lys	Thr	His	His	Asp	Met	Phe	Thr	Lys	Gly	Gln	Val	Thr	Thr	Gly	Gln
	290					295					300				
Tyr	Arg	Met	Leu	Ala	Lys	Arg	Gly	Gly	Tyr	Val	Trp	Val	Glu	Thr	Gln
305					310					315					320
Ala	Thr	Val	Ile	Tyr	Asn	Thr	Lys	Asn	Ser	Gln	Pro	Gln	Cys	Ile	Val
				325					330					335	
Cys	Val	Asn	Tyr	Val	Val	Ser	Gly	Ile	Ile	Gln	His	Asp	Leu	Ile	Phe

			340					345					350				
Ser	Leu	Gln	Gln	Thr	Glu	Cys	Val	Leu	Lys	Pro	Val	Glu	Ser	Ser	Asp		
		355					360					365					
Met	Lys	Met	Thr	Gln	Leu	Phe	Thr	Lys	Val	Glu	Ser	Glu	Asp	Thr	Ser		
	370					375					380						
Ser	Leu	Phe	Asp	Lys	Leu	Lys	Lys	Glu	Pro	Asp	Ala	Leu	Thr	Leu	Leu		
385					390					395					400		
Ala	Pro	Ala	Ala	Gly	Asp	Thr	Ile	Ile	Ser	Leu	Asp	Phe	Gly	Ser	Asn		
				405					410					415			
Asp	Thr	Glu	Thr	Asp	Asp	Gln	Gln	Leu	Glu	Glu	Val	Pro	Leu	Tyr	Asn		
			420					425					430				
Asp	Val	Met	Leu	Pro	Ser	Pro	Asn	Glu	Lys	Leu	Gln	Asn	Ile	Asn	Leu		
		435					440					445					
Ala	Met	Ser	Pro	Leu	Pro	Thr	Ala	Glu	Thr	Pro	Lys	Pro	Leu	Arg	Ser		
	450					455					460						
Ser	Ala	Asp	Pro	Ala	Leu	Asn	Gln	Glu	Val	Ala	Leu	Lys	Leu	Glu	Pro		
465					470					475					480		
Asn	Pro	Glu	Ser	Leu	Glu	Leu	Ser	Phe	Thr	Met	Pro	Gln	Ile	Gln	Asp		
				485					490					495			
Gln	Thr	Pro	Ser	Pro	Ser	Asp	Gly	Ser	Thr	Arg	Gln	Ser	Ser	Pro	Glu		
			500					505					510				
Pro	Asn	Ser	Pro	Ser	Glu	Tyr	Cys	Phe	Tyr	Val	Asp	Ser	Asp	Met	Val		
		515					520					525					
Asn	Glu	Phe	Lys	Leu	Glu	Leu	Val	Glu	Lys	Leu	Phe	Ala	Glu	Asp	Thr		
	530					535					540						
Glu	Ala	Lys	Asn	Pro	Phe	Ser	Thr	Gln	Asp	Thr	Asp	Leu	Asp	Leu	Glu		
545					550					555					560		
Met	Leu	Ala	Pro	Tyr	Ile	Pro	Met	Asp	Asp	Asp	Phe	Gln	Leu	Arg	Ser		
				565					570					575			
Phe	Asp	Gln	Leu	Ser	Pro	Leu	Glu	Ser	Ser	Ser	Ala	Ser	Pro	Glu	Ser		
			580					585					590				
Ala	Ser	Pro	Gln	Ser	Thr	Val	Thr	Val	Phe	Gln	Gln	Thr	Gln	Ile	Gln		
		595					600					605					
Glu	Pro	Thr	Ala	Asn	Ala	Thr	Thr	Thr	Thr	Ala	Thr	Thr	Asp	Glu	Leu		
	610					615					620						
Lys	Thr	Val	Thr	Lys	Asp	Arg	Met	Glu	Asp	Ile	Lys	Ile	Leu	Ile	Ala		

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<210> 331
<211> 92
<212> PRT
<213> Homo sapiens
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<400> 331
Met Ala Tyr Arg Gly Gln Gly Gln Lys Val Gln Lys Val Met Val Gln
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Pro Ile Asn Leu Ile Phe Arg Tyr Leu Gln Asn Arg Ser Arg Ile Gln
                20                      25                      30

Val Trp Leu Tyr Glu Gln Val Asn Met Arg Ile Glu Gly Cys Ile Ile
    35                      40                      45

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Gly Asn Asp Asn Asn Phe Val Ser Arg Glu Asp Cys Lys Arg Ala Cys
195 200 205

Ala Lys Ala Leu Lys Lys Lys Lys Lys Met Pro Lys Leu Arg Phe Ala
 210 215 220

Ser Arg Ile Arg Lys Ile Arg Lys Lys Gln Phe
 225 230 235

<210> 333
 <211> 291
 <212> PRT
 <213> Homo sapiens

<400> 333
 Met Gln Arg Ala Arg Pro Thr Leu Trp Ala Ala Ala Leu Thr Leu Leu
 5 10 15

Val Leu Leu Arg Gly Pro Pro Val Ala Arg Ala Gly Ala Ser Ser Gly
 20 25 30

Gly Leu Gly Pro Val Val Arg Cys Glu Pro Cys Asp Ala Arg Ala Leu
 35 40 45

Ala Gln Cys Ala Pro Pro Pro Ala Val Cys Ala Glu Leu Val Arg Glu
 50 55 60

Pro Gly Cys Gly Cys Cys Leu Thr Cys Ala Leu Ser Glu Gly Gln Pro
 65 70 75 80

Cys Gly Ile Tyr Thr Glu Arg Cys Gly Ser Gly Leu Arg Cys Gln Pro
 85 90 95

Ser Pro Asp Glu Ala Arg Pro Leu Gln Ala Leu Leu Asp Gly Arg Gly
 100 105 110

Leu Cys Val Asn Ala Ser Ala Val Ser Arg Leu Arg Ala Tyr Leu Leu
 115 120 125

Pro Ala Pro Pro Ala Pro Gly Asn Ala Ser Glu Ser Glu Glu Asp Arg
 130 135 140

Ser Ala Gly Ser Val Glu Ser Pro Ser Val Ser Ser Thr His Arg Val
 145 150 155 160

Ser Asp Pro Lys Phe His Pro Leu His Ser Lys Ile Ile Ile Ile Lys
 165 170 175

Lys Gly His Ala Lys Asp Ser Gln Arg Tyr Lys Val Asp Tyr Glu Ser
 180 185 190

Gln Ser Thr Asp Thr Gln Asn Phe Ser Ser Glu Ser Lys Arg Glu Thr
 195 200 205

Glu Tyr Gly Pro Cys Arg Arg Glu Met Glu Asp Thr Leu Asn His Leu
 210 215 220

005230" E35T3360

Lys Phe Leu Asn Val Leu Ser Pro Arg Gly Val His Ile Pro Asn Cys
225 230 235 240

Asp Lys Lys Gly Phe Tyr Lys Lys Lys Gln Cys Arg Pro Ser Lys Gly
245 250 255

Arg Lys Arg Gly Phe Cys Trp Cys Val Asp Lys Tyr Gly Gln Pro Leu
260 265 270

Pro Gly Tyr Thr Thr Lys Gly Lys Glu Asp Val His Cys Tyr Ser Met
275 280 285

Gln Ser Lys
290

<210> 334

<211> 582

<212> PRT

<213> Homo sapiens

<400> 334

Glu Ser Lys Gly Ala Ser Ser Cys Arg Leu Leu Phe Cys Leu Leu Ile
5 10 15

Ser Ala Thr Val Phe Arg Pro Gly Leu Gly Trp Tyr Thr Val Asn Ser
20 25 30

Ala Tyr Gly Asp Thr Ile Ile Ile Pro Cys Arg Leu Asp Val Pro Gln
35 40 45

Asn Leu Met Phe Gly Lys Trp Lys Tyr Glu Lys Pro Asp Gly Ser Pro
50 55 60

Val Phe Ile Ala Phe Arg Ser Ser Thr Lys Lys Ser Val Gln Tyr Asp
65 70 75 80

Asp Val Pro Glu Tyr Lys Asp Arg Leu Asn Leu Ser Glu Asn Tyr Thr
85 90 95

Leu Ser Ile Ser Asn Ala Arg Ile Ser Asp Glu Lys Arg Phe Val Cys
100 105 110

Met Leu Val Thr Glu Asp Asn Val Phe Glu Ala Pro Thr Ile Val Lys
115 120 125

Val Phe Lys Gln Pro Ser Lys Pro Glu Ile Val Ser Lys Ala Leu Phe
130 135 140

Leu Glu Thr Glu Gln Leu Lys Lys Leu Gly Asp Cys Ile Ser Glu Asp
145 150 155 160

Ser Tyr Pro Asp Gly Asn Ile Thr Trp Tyr Arg Asn Gly Lys Val Leu

	165		170		175										
His	Pro	Leu	Glu	Gly	Ala	Val	Val	Ile	Ile	Phe	Lys	Lys	Glu	Met	Asp
	180							185					190		
Pro	Val	Thr	Gln	Leu	Tyr	Thr	Met	Thr	Ser	Thr	Leu	Glu	Tyr	Lys	Thr
	195						200					205			
Thr	Lys	Ala	Asp	Ile	Gln	Met	Pro	Phe	Thr	Cys	Ser	Val	Thr	Tyr	Tyr
	210					215					220				
Gly	Pro	Ser	Gly	Gln	Lys	Thr	Ile	His	Ser	Glu	Gln	Ala	Val	Phe	Asp
225					230					235					240
Ile	Tyr	Tyr	Pro	Thr	Glu	Gln	Val	Thr	Ile	Gln	Val	Leu	Pro	Pro	Lys
				245					250						255
Asn	Ala	Ile	Lys	Glu	Gly	Asp	Asn	Ile	Thr	Leu	Lys	Cys	Leu	Gly	Asn
			260					265						270	
Gly	Asn	Pro	Pro	Pro	Glu	Glu	Phe	Leu	Phe	Tyr	Leu	Pro	Gly	Gln	Pro
	275						280					285			
Glu	Gly	Ile	Arg	Ser	Ser	Asn	Thr	Tyr	Thr	Leu	Thr	Asp	Val	Arg	Arg
	290					295					300				
Asn	Ala	Thr	Gly	Asp	Tyr	Lys	Cys	Ser	Leu	Ile	Asp	Lys	Lys	Ser	Met
305					310					315					320
Ile	Ala	Ser	Thr	Ala	Ile	Thr	Val	His	Tyr	Leu	Asp	Leu	Ser	Leu	Asn
				325					330					335	
Pro	Ser	Gly	Glu	Val	Thr	Arg	Gln	Ile	Gly	Asp	Ala	Leu	Pro	Val	Ser
			340					345					350		
Cys	Thr	Ile	Ser	Ala	Ser	Arg	Asn	Ala	Thr	Val	Val	Trp	Met	Lys	Asp
		355					360					365			
Asn	Ile	Arg	Leu	Arg	Ser	Ser	Pro	Ser	Phe	Ser	Ser	Leu	His	Tyr	Gln
	370					375					380				
Asp	Ala	Gly	Asn	Tyr	Val	Cys	Glu	Thr	Ala	Leu	Gln	Glu	Val	Glu	Gly
385					390					395					400
Leu	Lys	Lys	Arg	Glu	Ser	Leu	Thr	Leu	Ile	Val	Glu	Gly	Lys	Pro	Gln
				405					410					415	
Ile	Lys	Met	Thr	Lys	Lys	Thr	Asp	Pro	Ser	Gly	Leu	Ser	Lys	Thr	Ile
			420					425					430		
Ile	Cys	His	Val	Glu	Gly	Phe	Pro	Lys	Pro	Ala	Ile	Gln	Trp	Thr	Ile
	435						440					445			
Thr	Gly	Ser	Gly	Ser	Val	Ile	Asn	Gln	Thr	Glu	Glu	Ser	Pro	Tyr	Ile

450 455 460
 Asn Gly Arg Tyr Tyr Ser Lys Ile Ile Ile Ser Pro Glu Glu Asn Val
 465 470 475 480
 Thr Leu Thr Cys Thr Ala Glu Asn Gln Leu Glu Arg Thr Val Asn Ser
 485 490 495
 Leu Asn Val Ser Ala Ile Ser Ile Pro Glu His Asp Glu Ala Asp Glu
 500 505 510
 Ile Ser Asp Glu Asn Arg Glu Lys Val Asn Asp Gln Ala Lys Leu Ile
 515 520 525
 Val Gly Ile Val Val Gly Leu Leu Leu Ala Ala Leu Val Ala Gly Val
 530 535 540
 Val Tyr Trp Leu Tyr Met Lys Lys Ser Lys Thr Ala Ser Lys His Val
 545 550 555 560
 Asn Lys Asp Leu Gly Asn Met Glu Glu Asn Lys Lys Leu Glu Glu Asn
 565 570 575
 Asn His Lys Thr Glu Ala
 580

 <210> 335
 <211> 709
 <212> PRT
 <213> Homo sapiens

 <400> 335
 Met Ala Glu Val Glu Asp Gln Ala Ala Arg Asp Met Lys Arg Leu Glu
 5 10 15
 Glu Lys Asp Lys Glu Arg Lys Asn Val Lys Gly Ile Arg Asp Asp Ile
 20 25 30
 Glu Glu Glu Asp Asp Gln Glu Ala Tyr Phe Arg Tyr Met Ala Glu Asn
 35 40 45
 Pro Thr Ala Gly Val Val Gln Glu Glu Glu Glu Asp Asn Leu Glu Tyr
 50 55 60
 Asp Ser Asp Gly Asn Pro Ile Ala Pro Thr Lys Lys Ile Ile Asp Pro
 65 70 75 80
 Leu Pro Pro Ile Asp His Ser Glu Ile Asp Tyr Pro Pro Phe Glu Lys
 85 90 95
 Asn Phe Tyr Asn Glu His Glu Glu Ile Thr Asn Leu Thr Pro Gln Gln
 100 105 110

Leu	Ile	Asp	Leu	Arg	His	Lys	Leu	Asn	Leu	Arg	Val	Ser	Gly	Ala	Ala
115						120						125			
Pro	Pro	Arg	Pro	Gly	Ser	Ser	Phe	Ala	His	Phe	Gly	Phe	Asp	Glu	Gln
130						135						140			
Leu	Met	His	Gln	Ile	Arg	Lys	Ser	Glu	Tyr	Thr	Gln	Pro	Thr	Pro	Ile
145						150						155			
Gln	Cys	Gln	Gly	Val	Pro	Val	Ala	Leu	Ser	Gly	Arg	Asp	Met	Ile	Gly
			165						170			175			
Ile	Ala	Lys	Thr	Gly	Ser	Gly	Lys	Thr	Ala	Ala	Phe	Ile	Trp	Pro	Met
			180						185			190			
Leu	Ile	His	Ile	Met	Asp	Gln	Lys	Glu	Leu	Glu	Pro	Gly	Asp	Gly	Pro
195						200						205			
Ile	Ala	Val	Ile	Val	Cys	Pro	Thr	Arg	Glu	Leu	Cys	Gln	Gln	Ile	His
210						215						220			
Ala	Glu	Cys	Lys	Arg	Phe	Gly	Lys	Ala	Tyr	Asn	Leu	Arg	Ser	Val	Ala
225						230						235			
Val	Tyr	Gly	Gly	Gly	Ser	Met	Trp	Glu	Gln	Ala	Lys	Ala	Leu	Gln	Glu
			245						250			255			
Gly	Ala	Glu	Ile	Val	Val	Cys	Thr	Pro	Gly	Arg	Leu	Ile	Asp	His	Val
			260						265			270			
Lys	Lys	Lys	Ala	Thr	Asn	Leu	Gln	Arg	Val	Ser	Tyr	Leu	Val	Phe	Asp
275						280						285			
Glu	Ala	Asp	Arg	Met	Phe	Asp	Met	Gly	Phe	Glu	Tyr	Gln	Val	Arg	Ser
290						295						300			
Ile	Ala	Ser	His	Val	Arg	Pro	Asp	Arg	Gln	Thr	Leu	Leu	Phe	Ser	Ala
305						310						315			
Thr	Phe	Arg	Lys	Lys	Ile	Glu	Lys	Leu	Ala	Arg	Asp	Ile	Leu	Ile	Asp
			325						330			335			
Pro	Ile	Arg	Val	Val	Gln	Gly	Asp	Ile	Gly	Glu	Ala	Asn	Glu	Asp	Val
			340						345			350			
Thr	Gln	Ile	Val	Glu	Ile	Leu	His	Ser	Gly	Pro	Ser	Lys	Trp	Asn	Trp
355						360						365			
Leu	Thr	Arg	Arg	Leu	Val	Glu	Phe	Thr	Ser	Ser	Gly	Ser	Val	Leu	Leu
370						375						380			
Phe	Val	Thr	Lys	Lys	Ala	Asn	Ala	Glu	Glu	Leu	Ala	Asn	Asn	Leu	Lys
385						390						395			
												400			

Gln Glu Gly His Asn Leu Gly Leu Leu His Gly Asp Met Asp Gln Ser
 405 410 415
 Glu Arg Asn Lys Val Ile Ser Asp Phe Lys Lys Lys Asp Ile Pro Val
 420 425 430
 Leu Val Ala Thr Asp Val Ala Ala Arg Gly Leu Asp Ile Pro Ser Ile
 435 440 445
 Lys Thr Val Ile Asn Tyr Asp Val Ala Arg Asp Ile Asp Thr His Thr
 450 455 460
 His Arg Ile Gly Arg Thr Gly Arg Ala Gly Glu Lys Gly Val Ala Tyr
 465 470 475 480
 Thr Leu Leu Thr Pro Lys Asp Ser Asn Phe Ala Gly Asp Leu Val Arg
 485 490 495
 Asn Leu Glu Gly Ala Asn Gln His Val Ser Lys Glu Leu Leu Asp Leu
 500 505 510
 Ala Met Gln Asn Ala Trp Phe Arg Lys Ser Arg Phe Lys Gly Gly Lys
 515 520 525
 Gly Lys Lys Leu Asn Ile Gly Gly Gly Gly Leu Gly Tyr Arg Glu Arg
 530 535 540
 Pro Gly Leu Gly Ser Glu Asn Met Asp Arg Gly Asn Asn Asn Val Met
 545 550 555 560
 Ser Asn Tyr Glu Ala Tyr Lys Pro Ser Thr Gly Ala Met Gly Asp Arg
 565 570 575
 Leu Thr Ala Met Lys Ala Ala Phe Gln Ser Gln Tyr Lys Ser His Phe
 580 585 590
 Val Ala Ala Ser Leu Ser Asn Gln Lys Ala Gly Ser Ser Ala Ala Gly
 595 600 605
 Ala Ser Gly Trp Thr Ser Ala Gly Ser Leu Asn Ser Val Pro Thr Asn
 610 615 620
 Ser Ala Gln Gln Gly His Asn Ser Pro Asp Ser Pro Val Thr Ser Ala
 625 630 635 640
 Ala Lys Gly Ile Pro Gly Phe Gly Asn Thr Gly Asn Ile Ser Gly Ala
 645 650 655
 Pro Val Thr Tyr Pro Ser Ala Gly Ala Gln Gly Val Asn Asn Thr Ala
 660 665 670
 Ser Gly Asn Asn Ser Arg Glu Gly Thr Gly Gly Ser Asn Gly Lys Arg
 675 680 685

Ser Ser Tyr Glu Gln Asn Asp Asn Ser Leu Val Tyr Phe Ala Tyr Tyr
210 215 220

His Gly Leu Leu Gly Asn Arg Leu Trp Ser Ser Leu Gln Thr His Cys
 225 230 235 240
 Cys Ser Gln Asn Lys Cys Asn Phe Tyr Asp Asn Lys Asp Leu Glu Cys
 245 250 255
 Val Thr Asn Leu Gln Glu Val Ala Arg Ile Val Gly Asn Ser Gly Leu
 260 265 270
 Asn Ile Tyr Asn Leu Tyr Ala Pro Cys Ala Gly Gly Val Pro Ser His
 275 280 285
 Phe Arg Tyr Glu Lys Asp Thr Val Val Val Gln Asp Leu Gly Asn Ile
 290 295 300
 Phe Thr Arg Leu Pro Leu Lys Arg Met Trp His Gln Ala Leu Leu Arg
 305 310 315 320
 Ser Gly Asp Lys Val Arg Met Asp Pro Pro Cys Thr Asn Thr Thr Ala
 325 330 335
 Ala Ser Thr Tyr Leu Asn Asn Pro Tyr Val Arg Lys Ala Leu Asn Ile
 340 345 350
 Pro Glu Gln Leu Pro Gln Trp Asp Met Cys Asn Phe Leu Val Asn Leu
 355 360 365
 Gln Tyr Arg Arg Leu Tyr Arg Ser Met Asn Ser Gln Tyr Leu Lys Leu
 370 375 380
 Leu Ser Ser Gln Lys Tyr Gln Ile Leu Leu Tyr Asn Gly Asp Val Asp
 385 390 395 400
 Met Ala Cys Asn Phe Met Gly Asp Glu Trp Phe Val Asp Ser Leu Asn
 405 410 415
 Gln Lys Met Glu Val Gln Arg Arg Pro Trp Leu Val Lys Tyr Gly Asp
 420 425 430
 Ser Gly Glu Gln Ile Ala Gly Phe Val Lys Glu Phe Ser His Ile Ala
 435 440 445
 Phe Leu Thr Ile Lys Gly Ala Gly His Met Val Pro Thr Asp Lys Pro
 450 455 460
 Leu Ala Ala Phe Thr Met Phe Ser Arg Phe Leu Asn Lys Gln Pro Tyr
 465 470 475 480

<210> 337

<211> 543

<212> PRT

<213> Homo sapiens

275	280	285
Leu Ser Thr Ile Lys Ala Phe Ala Thr Gln Ser Gly Ser Gln Asp Leu		
290	295	300
Lys Ala Leu Asn Thr Ser Tyr Gln Ser Gln Leu Ile Lys Pro Ser Arg		
305	310	315 320
Met Arg Lys Tyr Pro Asn Arg Pro Ser Lys Thr Pro Pro His Glu Arg		
	325	330 335
Pro Tyr Ala Cys Pro Val Glu Ser Cys Asp Arg Arg Phe Ser Arg Ser		
	340	345 350
Asp Glu Leu Thr Arg His Ile Arg Ile His Thr Gly Gln Lys Pro Phe		
	355	360 365
Gln Cys Arg Ile Cys Met Arg Asn Phe Ser Arg Ser Asp His Leu Thr		
	370	375 380
Thr His Ile Arg Thr His Thr Gly Glu Lys Pro Phe Ala Cys Asp Ile		
385	390	395 400
Cys Gly Arg Lys Phe Ala Arg Ser Asp Glu Arg Lys Arg His Thr Lys		
	405	410 415
Ile His Leu Arg Gln Lys Asp Lys Lys Ala Asp Lys Ser Val Val Ala		
	420	425 430
Ser Ser Ala Thr Ser Ser Leu Ser Ser Tyr Pro Ser Pro Val Ala Thr		
	435	440 445
Ser Tyr Pro Ser Pro Val Thr Thr Ser Tyr Pro Ser Pro Ala Thr Thr		
	450	455 460
Ser Tyr Pro Ser Pro Val Pro Thr Ser Phe Ser Ser Pro Gly Ser Ser		
465	470	475 480
Thr Tyr Pro Ser Pro Val His Ser Gly Phe Pro Ser Pro Ser Val Ala		
	485	490 495
Thr Thr Tyr Ser Ser Val Pro Pro Ala Phe Pro Ala Gln Val Ser Ser		
	500	505 510
Phe Pro Ser Ser Ala Val Thr Asn Ser Phe Ser Ala Ser Thr Gly Leu		
	515	520 525
Ser Asp Met Thr Ala Thr Phe Ser Pro Arg Thr Ile Glu Ile Cys		
	530	535 540

<210> 338

<211> 148

<212> PRT

<400> 338

Leu Phe Leu Thr Ile Pro Phe Ala Phe Phe Leu Pro Glu Leu Ile Phe
20 25 30

Gly Phe Leu Val Trp Thr Met Val Ala Ala Thr His Ile Val Tyr Pro
35 40 45

Leu Leu Gln Gly Trp Val Met Tyr Val Ser Leu Thr Ser Phe Leu Ile
50 55 60

Ser Leu Met Phe Leu Leu Ser Tyr Leu Phe Gly Phe Tyr Lys Arg Phe
65 70 75 80

Glu Ser Trp Arg Val Leu Asp Ser Leu Tyr His Gly Thr Thr Gly Ile
85 90 95

Leu Tyr Met Ser Ala Ala Val Leu Gln Val His Ala Thr Ile Val Ser
100 105 110

Glu Lys Leu Leu Asp Pro Arg Ile Tyr Tyr Ile Asn Ser Ala Ala Ser
115 120 125

Phe Phe Ala Phe Ile Ala Thr Leu Leu Tyr Ile Leu His Ala Phe Ser
130 135 140

Ile Tyr Tyr His
145

<210> 339

<211> 196

<212> PRT

<213> Homo sapiens

<400> 339

Met Pro Gly Met Phe Phe Ser Ala Asn Pro Lys Glu Leu Lys Gly Thr

Thr His Ser Leu Leu Asp Asp Lys Met Gln Lys Arg Arg Pro Lys Thr
20 25 30

Phe Gly Met Asp Met Lys Ala Tyr Leu Arg Ser Met Ile Pro His Leu
35 40 45

Glu Ser Gly Met Lys Ser Ser Lys Ser Lys Asp Val Leu Ser Ala Ala
50 55 60

Glu Val Met Gln Trp Ser Gln Ser Leu Glu Lys Leu Leu Ala Asn Gln
65 70 75 80

Pro Gln Gly Phe Lys Ser Gly Asp Asp Leu Phe Pro Lys Asp Asp Lys

115 120 125
 Gly Asn Ala Ile Gly Gly Lys Ala Thr Phe Leu Asp Ala Trp Glu Ala
 130 135 140
 Met Glu Glu Leu Val Asp Glu Gly Leu Val Lys Ala Leu Gly Val Ser
 145 150 155 160
 Asn Phe Ser His Phe Gln Ile Glu Lys Leu Leu Asn Lys Pro Gly Leu
 165 170 175
 Lys Tyr Lys Pro Val Thr Asn Gln Val Glu Cys His Pro Tyr Leu Thr
 180 185 190
 Gln Glu Lys Leu Ile Gln Tyr Cys His Ser Lys Gly Ile Thr Val Thr
 195 200 205
 Ala Tyr Ser Pro Leu Gly Ser Pro Asp Arg Pro Trp Ala Lys Pro Glu
 210 215 220
 Asp Pro Ser Leu Leu Glu Asp Pro Lys Ile Lys Glu Ile Ala Ala Lys
 225 230 235 240
 His Lys Lys Thr Ala Ala Gln Val Leu Ile Arg Phe His Ile Gln Arg
 245 250 255
 Asn Val Ile Val Ile Pro Lys Ser Val Thr Pro Ala Arg Ile Val Glu
 260 265 270
 Asn Ile Gln Val Phe Asp Phe Lys Leu Ser Asp Glu Glu Met Ala Thr
 275 280 285
 Ile Leu Ser Phe Asn Arg Asn Trp Arg Ala Cys Asn Val Leu Gln Ser
 290 295 300
 Ser His Leu Glu Asp Tyr Pro Phe Asn Ala Glu Tyr
 305 310 315

<210> 341

<211> 422

<212> DNA

<213> Homo sapien

<220>

<221> misc_feature

<222> (1)...(422)

<223> n = A,T,C or G

<400> 341

gatganattt	ttncnagaga	gaggaagang	ctattcagtt	ggatgggatt	aatgcatca	60
caaataagag	aacttagaga	gaagtcggaa	aagtttgcct	tccaagcccg	aagttaacag	120
aatgatgaaa	cttatcatca	attcattgta	taaaaataaa	gagattttcc	tgagagaact	180
gatttcaa	atgcttgatg	ctttagataa	gataaggcta	atatcactga	ctgatgaaaa	240
tgctctttct	ggaaatgagg	aactaacagt	caaaattaag	tgtgataagg	agaagacctg	300

```
<210> 342
<211> 472
<212> DNA
<213> Homo sapien
```

```
<220>
<221> misc_feature
<222> (1) ... (472)
<223> n = A,T,C or G
```

```
<210> 343
<211> 139
<212> DNA
<213> Homo sapien
```

```
<210> 344
<211> 235
<212> DNA
<213> Homo sapien
```

```
<210> 345
<211> 458
<212> DNA
<213> Homo sapien
```

<400> 345						
ctgtaagg	ctattcag	ctgtgacc	tattttgg	tgtctttc	tactgttg	60
ctgttttg	acttcctg	aaaccgct	ctttgggt	gtgtcacc	gagctgtg	120
cataggac	cagttttg	ttaacct	aggcagtt	tatctcta	tttttcaa	180

```
<210> 346
<211> 525
<212> DNA
<213> Homo sapien
```

```
<220>
<221> misc_feature
<222> (1)...(525)
<223> n = A,T,C or G
```

```
<210> 347
<211> 423
<212> DNA
<213> Homo sapien
```

```
<210> 348
<211> 513
<212> DNA
<213> Homo sapien
```

<400>	348						
cctctaggcc	tgatgctctc	agaggcaata	gaagaaaagt	aaaaggaag	tctcacttca		60
cagacaatga	aaccctccta	accctcttcc	ccactaccca	caactcccta	cactgccaat		120
ctaaataaaa	agaggacaat	gcatgagtg	gagatacaca	tacacacaca	cacatacaca		180
cacacacacg	cacagcttcc	tttcagccaa	agaactgcaa	aatccttccc	cgggaaggagg		240
acaactggca	acaccaatca	aggcttgg	gtctaagg	atggctggaa	tcatgtgaga		300
ctggtataaaa	tccagggaga	aaatgtttca	ccttcagctc	attcccaagt	ctctatgaag		360

```

cccgccccac ttccacatag gggaaactgtg gctctggggg cagcctctgc agctactcag      420
aataggtggg aggaggggct ggctttgagg ctgccttagc catgaggctc tttgcctagg      480
aatagctgga gatgggagct gcagggggct cag                                     513

```

```

<210> 349
<211> 231
<212> DNA
<213> Homo sapien

```

```

<400> 349
ccttatttct cttgtccttt cgtacagggg ggaatttgaa gtagatagaa accgacctgg      60
attactccgg tctgaactca gatcacgtag gactttaatc gttgaacaaa cgaaccttta      120
atagcggtcg caccatcggt atgtcctgat ccaacatcga ggtcgtaaac cctattgttg      180
atatggactc tagagtagga ttgcgctggt atccctaggg taacttgttc c               231

```

```

<210> 350
<211> 341
<212> DNA
<213> Homo sapien

```

```

<400> 350
ctgcccaagg gcgttcgtaa cggaatgcc gaagcgtggg aaaaaggagg cgggtggcgga      60
agacggggat gagctcagga cagagccaga ggccaagaag agtaagacgg ccgcaaagaa      120
aatgacaaa gaggcagcag gagagggccc agccctgtat gaggaccccc cagatcagaa      180
aacctcacc agtggcaaac ctgccacacc caagatctgc tcttggaatg tggatgggct      240
tcgagcctgg attaagaaga aaggattaga ttgggttaaag gaagaagccc cagatatata      300
gtgccttcaa gagaccaaat gttcagagaa caaactacca g                    341

```

```

<210> 351
<211> 256
<212> DNA
<213> Homo sapien

```

```

<400> 351
ggcgttgggg acggtttagt gacgtggctc tttattcgtg agttttccat ttacctccgc      60
tgaacctaga gcttcagacg ccctatggcg tccgcctcga cccaaccggc ggccttgagc      120
gctgagcaag caaagggtgt cctcgcggag gtgatccagg cgttctccgc cccggagaat      180
gcagtgcgca tggacgaggc tcgggataac gcctgcaacg acatgggtaa gatgctgcaa      240
ttcgtgctgc ccgtgg                                     256

```

```

<210> 352
<211> 368
<212> DNA
<213> Homo sapien

```

```

<220>
<221> misc_feature
<222> (1)...(368)
<223> n = A,T,C or G

```

```

<400> 352
cctttcttgt aagtgaagaa naaggaatgc agcaaagaag agttcgacat tggagtcctt      60
agttccatca ggatcccatt cgcagccttt agcatcatgt agaagcaaac tgcacctatg      120
gctgagatag gtgcaatgac ctacaagatt ttgtgttttc tagctgtcca ggaaaagcca      180

```

tcttcagtct tgctgacagt caaagagcaa gtgaaacccat ttccagccta aactacataa 240
aagcagccga accaatgatt aaagacctct aagggtccat aatcatcatt aaatatgccc 300
aaactcattg tgacttttta ttttatatac aggattaaaa tcaacattaa atcatcttat 360
ttacatgg 368

<210> 353
<211> 368
<212> DNA
<213> Homo sapien

<400> 353
ctgagggtg gcagtaagca atgaggatgg gctataaagc tgttaactgg ctaagggcca 60
tccttgggca ggcatttcag acacatctgt agagagggca gtagcatctc cgataggcca 120
gctctgaagg aagcttaatg cttataacag tcacactgca taaattagct tagaatgctc 180
tcttgggtaa aaaatattaa tagtgtatat gcacttgaag agcaaaattc ctcaagaaaa 240
aaagtttaat agcaaggagt ttccatcagt cccgggtctt gtgaggatta ccacaacaaa 300
cacttaaaag gatacaacag gtacttatta aatgctgcct tgccttttac ctcttccttt 360
tttttttt 368

<210> 354
<211> 380
<212> DNA
<213> Homo sapien

<400> 354
ccatggcttc tcacccagac agtctttctg ggcaacttgg ggaagccctt gttctgctca 60
agtctcacc catggaagag gtgggggaag ggggccttgg tttttcagga agacagggtg 120
gagagcacga gtcactacaa agcagtaaaa gtgaatggtg tctccagggg ctgggtccag 180
aacaccacgg agagccccag ccataaagggt gtgttccgcc tctggcctgc aggaatctct 240
ttgaatctct ttgattggtg gctccaagag caatgggaag tcaacagcca ggaggctgga 300
ctgggttccc tgggaccccg aggtcccaga gctgctgggc agtggttgct ggcaaagaag 360
aaagggtcaa gagggtcagg 380

<210> 355
<211> 347
<212> DNA
<213> Homo sapien

<400> 355
ccagtggagg ggtgggggta tcatccccgc cgggggctgg cttggttget ggtgcctga 60
gcccttctct gccgcctgg gtgttgctt cactgatgga ggtaggcgtc cagccagatg 120
tcaccagact tcttcgggga cctgacgatg tccaccagcg cggtgaggaa gggttcaact 180
tcgtagctga ggccgtgctt ggcacacagc gacttgacca gcggggccac ccggctgtag 240
ttgtgtctcg gcatcctggg gaagaggtgg tgctcgatct ggaagttgag gtgcccgtg 300
aaccagttgg tgaaaagtga gggctccacg ttgcaggtgg ctgccag 347

<210> 356
<211> 157
<212> DNA
<213> Homo sapien

<400> 356
cctggagctg ctgaagactg ctattgggaa agctggctac actgataagg tggatcatcg 60
catggacgta gcggcctccg agttcttcag gtctgggaag tatgacctgg acttcaagtc 120

157

<400> 357

<400> 358

<400> 359

```
<210> 360
<211> 289
<212> DNA
<213> Homo sapien
```

<400> 360
 tttaaatttt actagtgtta cttaattgtat atttctaaaaa gagaatgcag taactaatgc 60
 cctaaatggt tgatctctgt ttgtcattac tttttcaaaa ttatTTTTTT ctgtaaagta 120
 taatatataa aacttcttgc ttaaattgaa tttctatatt agtggttaat tgcagtttat 180
 taaagggatc attatcagta atttcatagc aactgttcta gtgttttgtg tttttaaaac 240
 agaattagga atttgagata tctgattata tttttcatat gaatcacag 289

<210> 361
 <211> 311
 <212> DNA
 <213> Homo sapien

<400> 361
 ctgttcagta tggcaaaggg cagacttact ccttcatcca ctctgctgcc ttgatgaggt 60
 gaacacactg gaataagatg gagggcagga tacctgccaa agcctgagga atgagatgat 120
 ctgaaacaat tgggcaaagg ctggacattt caaaaagctg acttccaact gcagtttatg 180
 ggtatagaat ttgatgttcc cctcaagtcc tgactgctct ttctgaggca gccaggctag 240
 gccaagaaat gagctgctcc agcttctcca gagcacagca gcctcccagg gcctgtcagc 300
 atctgcagca g 311

<210> 362
 <211> 496
 <212> DNA
 <213> Homo sapien

<220>
 <221> misc_feature
 <222> (1)...(496)
 <223> n = A,T,C or G

<400> 362
 ccagtttcta aaanaatgca catttaaaga gaagcatcta ccacggcttt aaaacaaaac 60
 aactctgaga tgaacaatat gtgttatact cagagattaa caatctcaat catacatact 120
 gattctttca gacatttaac aaccactaca tttttttgca ttaatgaagt ttgactatat 180
 gtgtaaaggg actaaatatt tttgcaacag cctgttcttt gttcattctt ttctggatag 240
 cgtgtcctct gtattgcggt agatttatac attctgttgc ctaaatatgt gtgtaaaatg 300
 agctgataaa ctggagtact acttaaaaaa aagtctgtga ttataagat gcatatgctt 360
 tctatgtgaa tataagcttg tgcacaatgt ttaaaagaaa aacaatgaat tagaagagat 420
 cccccgtccc ccagtctgac atatttcata cagaatgttt aaaagaaaaa ctctgctagt 480
 cttggcaaac atttgg 496

<210> 363
 <211> 673
 <212> DNA
 <213> Homo sapien

<220>
 <221> misc_feature
 <222> (1)...(673)
 <223> n = A,T,C or G

<400> 363
 ccaagagggga gataanacaa acttctcaaa caaaaagaaa agaaaaacga atgattcatc 60
 tgctttaatc agtgtgatta atgcagcacc cattgccccg ggaaccgttt ctgctgtact 120

```

atctggatac taaaatgtta cggaagtagc tctttgttct ccctcactct gcccttagtt 180
aatagaaatt cagactcgcc aagtaaggct ttgtgcatag tgtcttcatt tcgcgtatag 240
ttgagcgctg tcttagcagt tggcttcatt gacagctcat tagtgttttg acttttctta 300
cccagcggtt attgaattct tgcttttaga caacttcctt tttgtagtgg tgaaccttgc 360
ccttttagtac agttcaagtg aatctggata attgttcatt tttgctttag cttagatacc 420
atgtagtggt ctgtggctac aggaagctgg ttctgtctgc ttccacagtc tgcttaaaaa 480
actgtctgac ttcgtgaata tagagaccaa gtttaccact tctgatgaag agaccaatta 540
agattcattc ctcatctgtt ttctttccag tgggagaaga gtcccatga aataagatga 600
aactgattcc atgcactagt acatgtaggc ttctcccttg cgcaaagctt aacaatttgt 660
aggaaacttt ggg 673

```

```

<210> 364
<211> 495
<212> DNA
<213> Homo sapien

<220>
<221> misc_feature
<222> (1)...(495)
<223> n = A,T,C or G

```

```

<400> 364
ccaaatgttt gcncaagact agcagagttt ttcttttaaa cattctgtat gaaatatgtc 60
agactggggg acgggggatc tcttctaatt cattgttttt cttttaaaca ttgtgcacaa 120
gcttatattc acatagaaag catatacatc ttataaatca cagacttttt ttttaagtagt 180
actccagttt atcagctcat tttacacaca tatttaggca acagaatgta taaatctacc 240
gcaatacaga ggacacacta tccagaaaag aatgaacaaa gaacaggctg ttgcaaaaat 300
atttagtccc tttacacata tagtcaaact tcattaatgc aaaaaatgta gtggttatta 360
aatgtctgaa agaatcagta tgtatgattg agattgttaa tctctgagta taacacatat 420
tgttcatctc agagttgttt tgttttaaag ccgtggtaga tgcttctctt taaatgtgca 480
tttttttagaa actgg 495

```

```

<210> 365
<211> 291
<212> DNA
<213> Homo sapien

```

```

<400> 365
aactgacaag cccttgcgcc tgctctcca ggatgtctac aaaattggtg gtattggtac 60
tgttcctgtt ggcccagtg gagactggtg ttctcaaacc cggtatggtg gtcacctttg 120
ctccagtcac cgttacaacg gaagtaaaat ctgtcgaaat gcacatgaa gctttgagtg 180
aagctcttcc tggggacaat gtgggcttca atgtcaagaa tgtgtctgtc aaggatgttc 240
gtcgtggcaa cgttgctggt gacagcaaaa atgacccacc aatggaagca g 291

```

```

<210> 366
<211> 277
<212> DNA
<213> Homo sapien

```

```

<400> 366
ctggatggtg cctcagaagg tgcattctgc ttctgcaggg gcttgaaaca ccaaggcact 60
ccagggatcc tggagtcaaa gcagcagccc cggttggtgc actccttggg ggtgacatgg 120
gggtagcccc cagtccaccc tgtccttggc tggcacggca cactggtttg cagacaggcc 180
cacgtactcc tcagcagagc tggaggacaa gcaaggccag gaccagcccc agcatgcaga 240

```

gcgctctggc agccatgacc accgtgggct ccgggac

277

<210> 367
<211> 311
<212> DNA
<213> Homo sapien

<400> 367
ccagagctgc ggggcctcag tacacggagc tgttccggat gccacagcac agcaccatgc 60
tcaggatcat ctggaagatc atgatcacag cgaccacgat ggcagcaatg ccgatgaggt 120
acagcttccc ggagaagagg tcatcgatct tctgggtggca gtcctccttg aagaggttgc 180
tgatgatgtt gctgcccagag ggacacaaat tgttcttgag cactgaggtg gtcaaagcag 240
tcagtgtgct ggagccacag cagtcaagcg tctcgtggaa ggtcttcacc acagccttgg 300
cgttgttggc g 311

<210> 368
<211> 384
<212> DNA
<213> Homo sapien

<400> 368
ccaaaggggt ctctagctgc tgcctctgctg ctccctgctca tggatgagtt tggcgatggg 60
gccggtgatg ccgcctatca aggtccagta ctcatcgaag ctgatgcgcc catcaggatt 120
ggcatccagg ttctggatga gcttatccgc agccttccgg ttccctgtgt ccgacagcat 180
gtggttcagc tctttctgga gcatctcgcg gaagctgctc ttgctgatct tgttcttgac 240
caggctgtac ctagacacat attttagtaa gttttccacc aggacaatga ctgccttctc 300
cagctccgtg tagcaagtct gacatctccc tgcttcgcct gctggcgggg cctaaggcgg 360
gggccaagcc cagttacagc ccag 384

<210> 369
<211> 216
<212> DNA
<213> Homo sapien

<400> 369
ccaagtgcc a ggtggctttc agcagcttcc tacgatcagc cgaagaaagc agaagctctg 60
gaggctgcc a tcgagaacct caatgaagcc aagaactatt ttgcaaagg t gactgcaaa 120
gagcgcac a gggacgtcgt ttacttccag gccagactct accataccct ggggaagacc 180
caggagagga accggtgtgc gatgctcttc cggcag 216

<210> 370
<211> 561
<212> DNA
<213> Homo sapien

<400> 370
ctggctcctt cttttgtggg cgtttggggg atgggctggt ttggggttta ggtgcagaga 60
atgggtttggg gccactgcgt actggaccac tctgagcctt cagggcaggg ttcttgtgag 120
tcttcatgtc atcagataca tgtttcaggg catgtgtaat gctctcccc tgattaatct 180
gcggaacag tctgagcgg gaagcagact catctgagcc tgaactggta gagactgggg 240
gaggaggggg gcctgggtgga gggggaggag gacctgatcc ggcagagggg ccagatggca 300
gtccgctcag ttcttttgcc acaggccccg ttttgcctca ggccagtcgg gtggtatgga 360
actccttaat gtaagcctgc agctctgtcc atatacttaa ataagctttg acccagtcta 420
catgcttctt atccacatct ttgtactctt tgaggactcg gtttgtataa aacatggcgg 480

```
<210> 371
<211> 518
<212> DNA
<213> Homo sapien
```

```
<210> 372
<211> 335
<212> DNA
<213> Homo sapien
```

```
<210> 373
<211> 467
<212> DNA
<213> Homo sapien
```

```
<210> 374
<211> 284
<212> DNA
<213> Homo sapien
```

<400> 374
tttccgtaaa agcgtgtaac aaggggtgtaa atatttataa ttttttatac ctgttgtgag 60

```
<210> 375
<211> 307
<212> DNA
<213> Homo sapien
```

```
<210> 376
<211> 650
<212> DNA
<213> Homo sapien
```

```
<220>  
<221> misc_feature  
<222> (1)...(650)  
<223> n = A,T,C or G
```

```
<210> 377
<211> 306
<212> DNA
<213> Homo sapien
```

```
<220>
<221> misc_feature
<222> (1)...(306)
<223> n = A,T,C or G
```

<400> 377
tctagatgca tgctcgagcg gccgccagtg tgatgganat ctgcagaatt cgccttctga 60
gcggccgccc gggcaggttc ggggtgctgcc ttcacctgcc aggcccttcc ccgctagctt 120

ggggcgagca gagctgcgtc cagtgggaact aaagccgttc caggattatc aaaaactgag 180
 cagcaacctt gggggacctg gatcatcacg gactcccca actggaaggc ccttctctgg 240
 cctcaattcc cgtctcaagg ccacgccttc cacctacagt ggagtcttcc gcaccagcg 300
 cgtcga 306

<210> 378
 <211> 199
 <212> DNA
 <213> Homo sapien

<220>
 <221> misc_feature
 <222> (1)...(199)
 <223> n = A,T,C or G

<400> 378
 ccacangtgg cacttgggtg tggtcctct gttatttgtc ctcatgtgag aaagcagatc 60
 atctccaaat cttgccattt gtatactttt ggtggagact tggatgtcat atcttctttg 120
 ttttgggttt tcttccttag cttattttgt ggcttttaaa gaagtggatt gtattgtgag 180
 atcctgtgat tcctgggtg 199

<210> 379
 <211> 216
 <212> DNA
 <213> Homo sapien

<220>
 <221> misc_feature
 <222> (1)...(216)
 <223> n = A,T,C or G

<400> 379
 ccagggcang tcatcaagag gggcattgtc ttgcatgcgg cctgccgtgt ccaccagcac 60
 caggtcaaag ctttggttac gtgcaaaagc aatggcttcc atggcaatgc cagcagcatc 120
 cttgccatag cccttttcaa acaactgcac catggtgcgg ccaccatgct tctctggagg 180
 gtgtagggca ctcaaagcc ggggtgtgtg acgcag 216

<210> 380
 <211> 555
 <212> DNA
 <213> Homo sapien

<400> 380
 ccatgggcct tcctttccac taaaaggaat tccgaacagc aaaaagaagg tcttgagata 60
 gtgaaaatgg tcatgatatc tttagaaggc gaagatgggt tggatgaaat ttattcattc 120
 agtgagagtc tgagaaaact gtgcgtcttc aagaaaattg agaggcattc cattcactgg 180
 ccctgccgac tgaccattgg ctccaatttg tctataagga ttgcagccta taaatcgatt 240
 ctacaggaga gagttaaaaa gacttggaca gttgtggatg caaaaaccct aaaaaagaa 300
 gatatacaaa aagaaacagt ttattgctta aatgatgatg atgaaactga agttttaaaa 360
 gaggatatta ttcaagggtt ccgctatgga agtgatatag ttcctttctc taaagtggat 420
 gaggaacaaa tgaaatataa atcggagggg aagtgttctc ctgttttggg attttgtaaa 480
 tcttctcagg gtcagagaag attcttcatg ggaaatcaag ttctaaaggc tttgccccaa 540
 gagatgatga ggcag 555

<210> 381
 <211> 406
 <212> DNA
 <213> Homo sapien

```

<400> 381
ctgcaccagg tgggcctcta ggtcccatta agcccattgg tccagggcca agtccaactc      60
cttttccatc atactgagca gcaaagttcc caccgagacc agggggggcca ggaggaccag      120
gtggaccagg agggcctgtg ggaccatctt caccatctct gcctgggggg cctgggtggac      180
ccctttctcc acgtggtcct ctatctccgg ctgggccctt tcttacagtt tcctcttgta      240
aagattggca tgttgctagg cataaggtta ctgcaagcag caacaaagtc cgcgtatcca      300
caaagctgag catgtctagc acttagacat gcagactcct tgtgtcgcag agcccctggg      360
tcaccggcgg aggtatcacc tggcggggcgc gggcatgcag tcgtgg                        406

```

<210> 382
 <211> 528
 <212> DNA
 <213> Homo sapien

<220>
 <221> misc_feature
 <222> (1)...(528)
 <223> n = A,T,C or G

```

<400> 382
ctgagcagtt tgtgggtntn tcttcccgcga agtttcagga agtattcaca aaagaaaaat      60
acattttttc cccagggggt ggggcaagga cagtggagag agtgctagga aatgagtccc      120
ctgggaaagg ggaccgggcc gtgatgttaa atatctccgg ctccaagtg actggatttg      180
cctaggacct tcagaccaac agacttcaga cctcagacc tgccccgggg ccagggtggag      240
aaagtgaggg cgtacaagg aagtgaatt ctgagttggt ggggctaagc ctgaccccct      300
ctccatgctc cccgccccaa ccaactctgg cctcagtaga tttttttttc agttgtggtt      360
gttgcccagg ctggagtgcg gtagcgccat cttggctcac tgcacctcca ccttcggggc      420
tcaagcgatt ctccagctc agcctcctga gtagctagga ctgcaggtgc tccaccacgc      480
ccggctaatt tttgtatttt tagtagagat ggggtttccc catgttgg                        528

```

<210> 383
 <211> 335
 <212> DNA
 <213> Homo sapien

<220>
 <221> misc_feature
 <222> (1)...(335)
 <223> n = A,T,C or G

```

<400> 383
ccatnttgag tctactcctg cgtcttgtgc cctagcaccc cgagaaccgt cagtttgagc      60
cagatggaag ctgagctgaa cacattacga tggatgatgg aaacataaga ctatcaagaa      120
atccaagtgg taatgggcga agtttattca gcatccggca atggacttat cgtagttggg      180
gaaacgggtg ttccgaataa tatcctggaa gttatcagga cacctatttt aaatataggc      240
ctgaattttg taaagtaata tttaagggtg tccgtgataa ttaaataaaa tgcttaattc      300
atgtggcgaa aaaaaaaaaa naaaaaaaaa aaaaaa                        335

```

<210> 384

<211> 333
 <212> DNA
 <213> Homo sapien

<400> 384
 agtccaatac ggctattggg gttgtagcag ctttcagagg aaattagtgg tctgggcttg 60
 cctccagctc cccaggggca gccccagtag ctacactgtc cagacagcac aagaccaggc 120
 tgggtgcacg tccatccgag cgctgectca gggatcgata aagtttcact gcagaaagtc 180
 tccactgcgg tatgctgaca tctgcccctga accttcaccc tacagcatta caggctttaa 240
 tcagattctg ctggaaagac acaggctgat ccacgtgacc tcttctgcct tcactgggct 300
 ggggtgatcc ttggtgcctt tgtttccaca agg 333

<210> 385
 <211> 343
 <212> DNA
 <213> Homo sapien

<400> 385
 ctgtgacacc tcaggttgaa agggctcttc tcttgaaca cccaccgagg ggcttgaggc 60
 aacagccagc cgatatggac ttctagctgc accgggtcac tgaggggtgga gaggtttgtc 120
 tggcacctgt actctccact gtcgtcgact gtggcagcgt caatgaagta gctcgaggcc 180
 tggcttgaga tgaggctctc attgtgaaac cactgtgtgg aattgtcctc aggggagtag 240
 gctccctggc acttcagagt cacactgtcc ttctcgagca cctgtacca ttgaggctcc 300
 aggaacacca cagccttttg gagatcttca gtccgcacgc caa 343

<210> 386
 <211> 244
 <212> DNA
 <213> Homo sapien

<400> 386
 tattctttga ttcttggcaa ataggtgaga gaactaatag caaccaggca actgaggacg 60
 aagtcaaaaa gtcggtaaca gaagaatgga atcagccaac ccacttgata agaaattgct 120
 ccataaacca gcattgaact gattataaac ataagaacag agacggcaaa aagaacacag 180
 gcattatcag ccattctctc agacgaatag taattaccga tgacttcata ctgaatgttg 240
 acag 244

<210> 387
 <211> 504
 <212> DNA
 <213> Homo sapien

<400> 387
 atctggagtc cagcctcagg gatgcgctac tttccattct ctgcattgaa cattogttct 60
 gtcagcatcc gctccagctt cactgcatca gcggcaaact tgcggatccc gtcagagagc 120
 ttctccacag ccattctggc ctcgttgtgc aaccaacgga aagacttctc atccagggtg 180
 attttttcca ggtcactggc ttgggccgcc ttggctgaga gcacaggcac cagcttggcg 240
 ttgtcctgca gcagctctcc caggagcttg gttgggatgg tgaggaagtc acagccggcc 300
 agtgctttga tctcgcccggt gttgcggaag gaggcgcccc tgacaatggg tttgtagcta 360
 aacttcttgt agtagttgta gatttttagtg acactcttta ccccagggtc ttccaggggc 420
 tcataggatt tcttgtcggg gtttgccaca tgccaatcaa ggatgcgccc aacaaatggg 480
 gagatgaggg tcacaccgc ctcg 504

<210> 388

```
<220>
<221> misc_feature
<222> (1)...(450)
<223> n = A,T,C or G
```

```
<210> 389
<211> 297
<212> DNA
<213> Homo sapien
```

```
<210> 390
<211> 223
<212> DNA
<213> Homo sapien
```

```
<210> 391
<211> 365
<212> DNA
<213> Homo sapien
```

<400> 391						
ctgaggaaga	aatgaaaaaa	gaccctgtcc	ctcatggccc	gcccaactggc	ctcctgtgaa	60
ctctgtcctg	ttgccaaacc	cagatgaagt	cagccaaaaa	gtgctttcca	catcctctct	120
ctggggctgc	ccagcctgac	cgtaggggat	ccactggcag	agccaagggtg	gatgctgggtg	180
cctgaagctg	gaagccagca	ggacatgaga	ccctcctgt	agcaggaagt	ggttctagaa	240
ctcccagcag	aacagaacgg	aaaaggagct	gattggggat	agaatgagtt	ctgctaaaca	300
qccagatqct	ctgaqagagg	tgacactgga	ctgtctcgga	ggtgtgtgca	gatggctaca	360

365

<400> 392

```
<210> 393
<211> 213
<212> DNA
<213> Homo sapien
```

```
<220>
<221> misc_feature
<222> (1)...(213)
<223> n = A,T,C or G
```

<400> 393

ccaataatca	agnacaaana	ctggatttga	ggatggatca	gttctgaaac	agtttctttc	60
tgaacagag	aaaatgtccc	ctgaagacag	agcaaaatgc	tttggaaga	atgaggccat	120
acaggcagcc	catgatgocg	tggcacagga	aggccaatgt	cgggtagatg	acaagggtgaa	180
tttccatttt	attctgttta	acaacgtgga	tgg			213

```
<210> 394
<211> 334
<212> DNA
<213> Homo sapien
```

<400> 394

cctaccata	atccagagag	gcttgcccag	aggaggacta	cgtgggggac	gtgccaccag	60
aaccctactt	gggggcggga	tgtcactccg	aggtcaaaac	ctgctccgag	gtggacgagc	120
cgtagctccc	cgaatgggct	taagaagagg	tggtgttcga	ggtcgtggag	gtcctggggag	180
agggggccta	gggcgtggag	ctatgggtcg	tggcggaaatc	ggtggtagag	gtcgggggtat	240
gataggctcg	ggaagagggg	gctttggagg	cagaggccga	ggcctggaac	gagggagagg	300
tgccttcgtg	gcgcctgtat	tgaaccaagga	cgag			334

```
<210> 395
<211> 174
<212> DNA
<213> Homo sapien
```

```
<210> 396
<211> 140
<212> DNA
<213> Homo sapien
```

```
<210> 397
<211> 318
<212> DNA
<213> Homo sapien
```

```
<210> 398
<211> 517
<212> DNA
<213> Homo sapien
```

<400> 398						
ccttncttcg	ccatccattc	atcgaccctc	tccagcactt	gctgcaggct	tggctgacca	60
tccaccatgg	cttgaataat	ccgggtgagc	tctgtacaga	atggggtaag	ctgtggatgg	120
actacaggct	ggacatacat	gtgaaaggta	gactcaatct	ccatgggtccg	gccatttagc	180
tttaggatgg	ggaactcgat	gatttcttga	ggatgaatct	gtggcttgct	gcacgtggcc	240
tcaaagtcca	gcactaaaaa	gtagtgtatac	ctctggagag	ggaaggacac	cattgccgcc	300
atggatgcgc	caaagccgtg	ggccgccagc	tttctggttg	atatggagca	gaactccgga	360
acaccacagg	gagaaaataa	gtgggagccc	agcacttttc	ttgctcttga	aagtaaatac	420
gaagaaaatc	gagctgctcc	agtctgtaaa	ggtgctagca	ttgaacatcc	agaagcatct	480
aaaactctcc	ttacttcgaa	gatgccaaaga	ccggcag			517

<210> 399
 <211> 329
 <212> DNA
 <213> Homo sapien

<400> 399
 ccaacctcag gcaacgggtg gagcagtttg ccagggcctt ccccatgcct ggttttgatg 60
 agcattgaag gcacctggga aatgaggccc acagactcaa agttactctc ctcccccta 120
 cctggggcag tgaaatagaa agcctttcta ttttttggtg cgggagggaa gacctctcac 180
 ttagggaag agccaggtat agtctccctt cccagaatth gtaactgaga agatcttttc 240
 tttttccttt tttcgtaac aagacttaga aggagggccc aggcactttc tgtttgaacc 300
 cctgtcatga tcacagtgtc agagacgcg 329

<210> 400
 <211> 451
 <212> DNA
 <213> Homo sapien

<400> 400
 ctggcttcac tgctcaggtg attatcctga accatccagg ccaaataagc gccggctatg 60
 cccctgtatt ggattgccac acggctcaca ttgcatgcaa gtttgctgag ctgaaggaaa 120
 agattgatcg ccgttctggt aaaaagctgg aagatggccc taaattcttg aagtctggtg 180
 atgctgccat tgttgatatg gttcctggca agcccatgtg tgttgagagc ttctcagact 240
 atccaccttt gggtcgcttt gctgttcgtg atatgagaca gacagttgag gtgggtgtca 300
 tcaaagcagt ggacaagaag ctgctggagc tggcaaggct accaagtctg cccagaaagc 360
 tcagaagcta aatgaatatt atccctaata cctgccaccc cactcttaat cagtgggtgga 420
 agaacggctc agaactgttt gtttcaattg g 451

<210> 401
 <211> 180
 <212> DNA
 <213> Homo sapien

<400> 401
 ccaggaagca ggccagggga ttggcagcac tgcccagcac cacagccagg ttgtaggcca 60
 gacgcccgtg gggtaagcag gaaaagctct gcacggcagg cagcacgcca ttggtcagcg 120
 cgttgggtggc ggccaacagg cccagcaggc aggcactgag ggctgataga agctgatagg 180

<210> 402
 <211> 385
 <212> DNA
 <213> Homo sapien

<400> 402
 ccaggccacc tgtgcggggc tcctcgatgt ggaaggttcg ggtgaggaga ttgtagaagg 60
 agccgtagca cacggccacc acagtgcacg tgaggcagat cacgttgtag ggcatgctga 120
 agtccggtgt cggcaggttc accagcagcg gctccgtgta gagccgcaca aagtagttag 180
 agccatcaga gactgggaac aggctgttga agaggggact ctcttcccag tccactggct 240
 tggctgctac catgctgggc acaagggcgc tgaggacaga tgggctgaca tagaagccat 300
 ggtaggtatc tggcgtgtac tgggtccact tcagcagcgc ccgctcaaac tggatggaaa 360
 ccttggtgac tgagttggcc ggcag 385

<210> 403
 <211> 440

<212> DNA
<213> Homo sapien

<220>
<221> misc_feature
<222> (1)...(440)
<223> n = A,T,C or G

<400> 403
ctgtttaacc agnaacccgg ggggtcaccc cccacagaat gtacatgaaa cactagagga 60
ctgcatgttt ttccctgaga gaagcgtaag acaaacagaa gtcaaaaagt agtcactggg 120
agcgccatcc ttctaagcaa atcctccctt tcccttttgg aggatttgcc cgaactacgt 180
agccagtcag cacttagacc acctgcctcc tccccccct ataaaccac cactcccctc 240
ctcctttccc aaaccacttg ggggtgccta agccctcact gcccgaagcc caaaatatca 300
gctaagatcc ttgtcagtat ttccacagtc atacctaatg aattgggaag tggggcccct 360
aaaaaccaat tcacatctat gcacttggtt ccactggatt tggcagacag gcttttttag 420
ttaccgtaac cagatcttaa 440

<210> 404
<211> 239
<212> DNA
<213> Homo sapien

<400> 404
cctacgaaaa actcccggcc ggtgaagaga acgtcagtgc catccagcgt cgcgttctcg 60
tctcctatct ccacaattcg gagccccagg tcttgacagg ctttgoggac tccatcgacc 120
tctggcctac gagcggggct ccagggccgc gtgattaggg ccgtgtcccc ttggatcacg 180
gccgtgtcgc caagcagcgg tcccagcggc aatgactcct cagggtggcag ttctagcag 239

<210> 405
<211> 261
<212> DNA
<213> Homo sapien

<400> 405
ctggagaggc agcccttcac cggatgccc a gctccgtgcc cctgcggggc ccagcacagt 60
ttaccttctc cccccacggc ggtcccatct actctgtgag ctgttcccc ttccacagga 120
atctcttctc gagcgtctgg actgacgggc atgtccacct gtactccatg ctgcaggccc 180
ctcccttgac ttcgctgcag ctctccctca agtatctgtt tgctgtgcgc tgggtccccag 240
tgcgccctt ggtttttgca g 261

<210> 406
<211> 641
<212> DNA
<213> Homo sapien

<220>
<221> misc_feature
<222> (1)...(641)
<223> n = A,T,C or G

<400> 406
ctgctcccg gcntgggtggc agcaagtaga catcgggcct gtgcagggcc acccccttgg 60
gccgggagat ggtctgcttc agtggcgagg gcaggctctgt gtgggtcacg gtgcacgtga 120

```
<210> 407
<211> 173
<212> DNA
<213> Homo sapien
```

```
<210> 408
<211> 165
<212> DNA
<213> Homo sapien
```

```
<210> 409
<211> 329
<212> DNA
<213> Homo sapien
```

```
<210> 410
<211> 235
<212> DNA
<213> Homo sapien
```

<400> 410

```

ccatcagnga gaaaggtggt tgtcagttgt ttcacaaacc agattgagga ggacaaactg      60
ctctgccaat ttctggattt ctttattttt agcaaacact ttctttaaag cttgactgtg      120
tgggcactca tccaagtgat gaataatcat caagggtttg ttgcttgtct tggatttata      180
tagagctttt tcatatgtct gagtccagat gagttggtca cccaacctc tggag          235

```

```

<210> 411
<211> 294
<212> DNA
<213> Homo sapien

```

```

<400> 411
aattaaggga agatgaagat gataaaacag ttttggatct tgctgtggtt ttgtttgaaa      60
cagcaacgct tcggtcaggg tatcttttac cagacactaa agcatatgga gatagaatag      120
aaagaatgct tcgcctcagt ttgaacattg accctgatgc aaaggtggaa gaagagcctg      180
aagaagaacc tgaagagaca gcagaagaca caacagaaga cacagagcaa gacgaagatg      240
aagaaatgga tgtgggaaca gatgaagaag aagaaacagc aaaggaatct acag          294

```

```

<210> 412
<211> 433
<212> DNA
<213> Homo sapien

```

```

<220>
<221> misc_feature
<222> (1)...(433)
<223> n = A,T,C or G

```

```

<400> 412
cctgagaagc cagaggcagg tggagagggg gtggaaagtg agcagcgggc tgggctggag      60
ccgcacacgc tctcctccca tgtaaataag cacctttaga aaaattcaca agtccccatc      120
cacaaaaaaa aaaanaanaa aaatttcagg gantaaaaat anactttgaa caaaaaggaa      180
catttgntgg cctggggggg catctnantt tntntagcnc cagngattcc ctccccnccc      240
cacccatcac atanatgtaa cacctttggt ntaaaatggg gagccgtttc caccntgccc      300
ccntccccgc cccagggcag ttgccccggn gacacntcaa gacaggancg aggtagtntt      360
tcancancac agttncacaa ggaacagaac agtntctccc gccagccct gcggcacaag      420
ggattgacac gcn          433

```

```

<210> 413
<211> 494
<212> DNA
<213> Homo sapien

```

```

<220>
<221> misc_feature
<222> (1)...(494)
<223> n = A,T,C or G

```

```

<400> 413
ccttatttct cttgtcnctt cgtacagggg ggaatttgaa gtagatagaa accgacctgg      60
attactccgg tctgaactca gatcacgtag gactttaatc gttgaacaaa cgaaccttta      120
atagcggctg caccatcggg atgtcctgat ccaacatcga ggtcgtaaac cctattgttg      180
atatggactc tagaatagga ttgcgctggt atccctaggg taacttgttc cgttggtcaa      240
gttattggat caattgagta tagtagttcg ctttgactgg tgaagtctta gcatgtactg      300
ctcggagggt gggttctgct ccgaggtcgc cccaaccgaa atttttaatg cagggttggt      360

```

```

agtttaggac ctgtgggttt gttagggtact gtttgcatta ataaattaaa gtcctatagg 420
gtcttctcgt cttgctgtgt tatgcccgcc tcttcacggg cagggtcaatt tcaactggta 480
aaagtaagag acag 494

```

```

<210> 414
<211> 294
<212> DNA
<213> Homo sapien

```

```

<400> 414
ctgggcggat agcaccgggc atatttttga atggatgagg tctggcaccg tgagcagtc 60
agcgaggact tggctcttagt tgagcaattt ggctaggagg atagtatgca gcacggttct 120
gagtctgtgg gatagctgcc atgaagtaac ctgaaggagg tgctggctgg taggggttga 180
ttacaggggt gggaacagct cgtacacctg ccattctctg catatactgg ttagtgaggt 240
gagcctggcg ctcttctttg cgtgagcta aagctacata caatggcctt gtgg 294

```

```

<210> 415
<211> 421
<212> DNA
<213> Homo sapien

```

```

<400> 415
ccttgccccct gccctccac gaatgggttaa tatatatgta gatatatatt ttagcagtga 60
cattcccaga gagccccaga gctctcaagc tcctttctgt cagggtgggg ggttcagcct 120
gtcctgtcac ctctgagggt cctgctggca tcctctcccc catgcttact aatacattcc 180
cttccccata gccatcaaaa ctggaccaac tggcctcttc ctttccccctg ggacccaaat 240
ttaggggcct cagtcctca ccgccatgcc ctggcctatt ctgtctctcc ttcttcccc 300
tggcctgttc tgtctctgag ctctgtgtcc tccgttcatt ccatggctgg gagtcaactga 360
tgctgcctct gccttctgat gctggactgg ccttgcttct acaagtatgc ttctcccaca 420
g 421

```

```

<210> 416
<211> 342
<212> DNA
<213> Homo sapien

```

```

<220>
<221> misc_feature
<222> (1) ... (342)
<223> n = A,T,C or G

```

```

<400> 416
ccactttctt tcccacnctg gaaggcggca tctatgactt cattggggag ttcatgaagg 60
ccagcgtgga tgtggcagac ctgataggtc taaaccttgt catgtcccgg aatgccggca 120
agggagagta caagatcatg gttgctgccc tgggctgggc cactgctgag cttattatgt 180
cccgtgcac tcccctatgg gtcggagccc ggggcattga gtttgactgg aagtacatcc 240
agatgagcat agactccaac atcagtctgg tccattacat cgtcgcgtct gctcaggtct 300
ggatgataac acgctatgat ctgtaccaca ctttccggcc gg 342

```

```

<210> 417
<211> 389
<212> DNA
<213> Homo sapien

```

<400> 417

tattaattag	gttcttaaga	catttagaac	accaatttgt	gaggataaat	tccattcgtc	60
agagcaaaca	cagatcgcag	gtagccctgg	agctgaggaa	tagctttgat	ttttggtaaa	120
at ttgtgagt	ccacagcttt	ctgatcaatc	ttgcgctgct	ccgtaatctc	atatttctct	180
ttttctgtgt	cgaagatctc	accttcctgg	tgtctgggct	tccgcagctt	cttcttcttg	240
aagtaagcat	cagtaagatg	ttttgggatt	tttacattgc	tgatatcgat	tttggttgaa	300
gtggcaatga	caaatttctg	gtgtgttctt	cgtagaggaa	ctcgattgag	gaccagaggt	360
ccagtcacaa	gtaataagcc	actagccag				389

<210> 418

<211> 343

<212> DNA

<213> Homo sapien

<400> 418

gtgggagggga	gccaggttgg	gatggagggga	gtttacagga	agcagacagg	gccaacgtcg	60
aagccgaatt	cctggctctgg	ggcaccaacg	tccaaggggg	ccacatcgat	gatgggcagg	120
cgggaggtct	tggtggtttt	gtattcaatc	actgtcttgc	cccaggctcc	gggtgtgactc	180
gtgcagccat	cgacagtgc	gctgtagggtg	aagcggctgt	tgccctcggc	gcggatctcg	240
atctcgttgg	agccctggag	gagcagggcc	ttcttgaggt	tgccagtctg	ctggtccatg	300
taggccacgc	tgtttttgca	gtggtagggtg	atgttctggg	agg		343

<210> 419

<211> 255

<212> DNA

<213> Homo sapien

<400> 419

cctagcaaga	gaatcaccaa	at ttatggag	agttaacagg	ggtttaacag	gaaggaagtg	60
ccttttagtaa	gttctcaagc	cagaggctgg	aggcagcagc	taaatcagag	gacagcatcc	120
tcagtgaag	tgagccattc	ggggtggcat	gtcactccag	gaataaacac	aacttagaaa	180
caaattgattt	cgtaggatag	cacagtgcac	tggtgcactg	tgaacctgag	gccactgtgt	240
caaactgtgc	actgg					255

<210> 420

<211> 261

<212> DNA

<213> Homo sapien

<400> 420

cttctgatga	taaccaaccc	ctagctacca	ctctgtattc	atcaggggag	gggtataaac	60
cccacatgca	agaagaaccc	ttgccccag	tgtcaaatgg	gatggggatg	ctagagttat	120
agtaaagggg	aaaccctatg	taagctgtta	acagagttca	caggggtagg	gataaccct	180
gttctccagc	tcccaaattg	gctcactttc	ccagcttctt	catccgttca	tcaatgctgg	240
caaagttccc	ctcaactgtg	g				261

<210> 421

<211> 179

<212> DNA

<213> Homo sapien

<400> 421

ccttcctgtt	gttgtttcaa	atgctgcttg	atttctcgta	acagatctgc	atctatgtaa	60
tacctttctt	cagatctgac	tgctccaaaa	tgattctgca	tcttgatttg	agacatcaat	120

tcatttagtc ggcccttgaa ctgagtaggt gcatttagtt caccctgaat cgtatccag 179

<210> 422
 <211> 424
 <212> DNA
 <213> Homo sapien

<400> 422
 cgaggtccaa atctgatctg cagatgcaga agattcgaca gaagctgcag actaaacagg 60
 ctgccatgga gaggtctgga aaagctaagc aactgcgagc acttaggaaa tacgggaaga 120
 aggtgcaaac ggaggttctt cagaagaggc agcaggagaa agcccatatg atgaatgcta 180
 ttaagaaata tcagaaaggc ttctctgata aactggatth ccttgaggga gatcagaaac 240
 ctctggcaca gcacaagaag gcaggagcca aaggccagca gatgaggaag gggcccagtg 300
 ctaaacgacg gtataaaaac cagaagtttg gttttggtgg aaagaagaaa ggctcaaagt 360
 ggaacactcg ggagagctat gatgatgtat ctagcttccg ggccaagaca gctcatggca 420
 gagg 424

<210> 423
 <211> 256
 <212> DNA
 <213> Homo sapien

<400> 423
 ctgtggccta gggctacctc aagactcacc tcattccttac cgcacattta aggcgccatt 60
 gcttttggga gactggaaaa gggaagggtg ctgaaggctg tcaggattct tcaaggagaa 120
 tgaatactgg gaatcaagac aagactatac cttatccata ggcgaggtg cacaggggga 180
 ggccataaag atcaaactg catggatggg tcctcacgca gacacacca cagaaggaca 240
 ctagcctgtg cacgcg 256

<210> 424
 <211> 330
 <212> DNA
 <213> Homo sapien

<400> 424
 ccagccgcat gggagtggag gcagtcacg ccttgctaga ggccaccccg gacacccag 60
 cttgcgtcgt gtcactgaac gggaaccacg ccgtgcgcct gccgctgatg gaggcgtgc 120
 agatgactca ggatgtgcag aaggcgatgg acgagaggag atttcaagat gcggttcgac 180
 tccgagggag gagctttgag ggcaacctga acacctaca gcgacttgcc atcaagctgc 240
 cggatgatca gatcccaaag accaattgca acgtagctgt catcaacgtg ggggcacccg 300
 cggctgggat gaacgcggcc gtacgctcag 330

<210> 425
 <211> 333
 <212> DNA
 <213> Homo sapien

<220>
 <221> misc_feature
 <222> (1)...(333)
 <223> n = A,T,C or G

<400> 425
 ctgctccatg gnetcaaagt cagcaccacc cacaccaca atgatcactg acatgggcag 60

```

gttcgaggca cgcaccacag cctcacgtgt ggcttccaca tccgtcacag caccatcagt 120
cagnagaaac agnatgaagt attgngaggg antccctga tgtgcagcct gggctgcaaa 180
cctggacctg cccgggaggc cgctcgaaaag ggcgaattcc agcacactgg cggccgttac 240
tagnggatnc aganctcggg acnaagcttg gcagtaatca tggtcatagc tgtttcctgt 300
gagcggttg gatgaacgag gccgtacgt cat 333

```

```

<210> 426
<211> 411
<212> DNA
<213> Homo sapien

```

```

<220>
<221> misc_feature
<222> (1)...(411)
<223> n = A,T,C or G

```

```

<400> 426
gggtgttcat catgaggatt gcttctgcc a tggagctgat ggacgtgggc aggttgctga 60
gaaggtgggg tggaagtga tgccgggggt ggggtgagtgc cctggctctg ttcataagggg 120
agcctttccc tagcagtga acgctgtggg cattttctct agcatattcc cttgggaagt 180
ctagatttgc tattaatctg gctgagaatc taagttctgt gccttagaga cagtttgac 240
tttcccatat tgtgcctggg acagccatat gatttttttt cccaccaaac aagtatgcaa 300
acagaaacca gttcaaagg ggatgggtga aaagatgagg cagtanaaat gcctttgaat 360
ggttttctgt agctaattct ctttaaattt tgtcctgctt tttttcttta t 411

```

```

<210> 427
<211> 450
<212> DNA
<213> Homo sapien

```

```

<220>
<221> misc_feature
<222> (1)...(450)
<223> n = A,T,C or G

```

```

<400> 427
acgtgtacaa gtttgaactg gatacctctg aaagaaagat tgaatttgac tctgcctctg 60
gcacctacac tctctactta atcattggag atgccacttt gaagaaccca atcctctgga 120
atgtggctga tgtggnatc aagttccctg aggaagaagc tccctcgact gtcttgccc 180
agaacctttt cactccaaaa caggaaattc agcacctgtt ccgcgagcct gagaagaggc 240
ccccaccgt ggtgtccaat acattcactg ccctgatcct ctgcgcgttg cttctgctct 300
tcgctctgtg gatccggatt ggtgccaatg tctccaactt cacttttgct cctagcacga 360
ttatatttca cctgggacat gctgctatgc tgggactcat gtatgtctac tggactcagc 420
tcaacatgtt ccagacctg aagtacctg 450

```

```

<210> 428
<211> 377
<212> DNA
<213> Homo sapien

```

```

<220>
<221> misc_feature
<222> (1)...(377)
<223> n = A,T,C or G

```

<400> 428

```

cagggctata gtgcgctatg ttgatctggt gttcatgcta agttccgcat caatatggtg      60
acttcttggg agtggggggac caccagggttg cctaaggagg ggtgaacctg cctacgttgg      120
aaatagagct ggncaaaaact cctgtgctca tcagtagtag aattgcacct gtgaatagcc      180
nccgccctcc agcatgggca acataacaag accctgcctc ttaaagataa aaattggaaa      240
acactngtag gaaaaaaagg gtgnnttggtc taaataaatn tggattgggn ataaatgacn      300
caaaactatc atgaatttga aagcntttct aatttcttga aagtctgaaa aaagttaaan      360
cncaatttta tctnaaa                                     377

```

<210> 429

<211> 206

<212> DNA

<213> Homo sapien

<400> 429

```

gttgctcctc caaagaaggt tggcttcaag gccgtgtcca gggacccacg agcagaggca      60
ctgggggggca agggatctcc aaggggggcaa gggatcccta aagggggtag ctcacagggtg      120
aggggggttta gggccctct agggagcgcc tgaggccata cattcaagag tgtccctggg      180
gaggcccgag gaagagccag gactgg                                     206

```

<210> 430

<211> 473

<212> DNA

<213> Homo sapien

<220>

<221> misc_feature

<222> (1)...(473)

<223> n = A,T,C or G

<400> 430

```

ccttatttnt cttgtccttt cgtacaggga ggaatttgaa gtagatagaa accgacctgg      60
attactccgg tctgaactca gatcacgtag gactttaatc gttgaacaaa cgaaccttta      120
atagcggctg caccatcggg atgtcctgat ccaacatcga ggtcgtaaac cctattgttg      180
atatggactc tagaatagga ttgcgctggt atccctaggg taacttggtc cgttgggtcaa      240
gttattggat caattgagta tagtagttcg ctttgactgg tgaagtotta gcatgtactg      300
ctcggagggt gggttctgct ccgaggtcnc cccanccgaa atttttaatg cagggttggg      360
agntnaggac ctgtgggttt gttaggtact ggggtgcatta ataaattaaa gctccatagg      420
gtcttctcgt cttgctgtgt tatgccncc tcttcacggg caggtcaatt tca                                     473

```

<210> 431

<211> 215

<212> DNA

<213> Homo sapien

<220>

<221> misc_feature

<222> (1)...(215)

<223> n = A,T,C or G

<400> 431

```

cctgtatnaa gctanaaaaa gactaccagc ccgggatcac cttcatcgtg gtgcagaaga      60
ggcaccacac ccggctcttc tgcactgaca agaacgagcg ggttgggaaa agtggaaaca      120

```

ttccagcagg cacgactgtg gacacgaaaa tcacccaccc caccgagttc gacttctacc 180
tgtgtagtca cgctggcatc caggggacaa gcagg 215

<210> 432
<211> 391
<212> DNA
<213> Homo sapien

<220>
<221> misc_feature
<222> (1)...(391)
<223> n = A,T,C or G

<400> 432
ccagcactgc cacaaacttt ttcagggcca ccaggcgtg cccttcagg accgggaacc 60
tgcccacttc tatccgcagg atgtagtgc gtgcagattc caggtcagcc atgtagatcc 120
tggagcgatc tgccaatttc caaacagtgg gagctatctt gttagcagt gttgggtgcaa 180
ctgtggtctg ggcagcctcc ctggtgagcc cagagagtct ctgcaggtaa gcggtataga 240
aggacctgga ttccatgagc acggggactc gggagacgga gccattccgg aacagcaggt 300
agcaagaggg gaagtcggtg acaccaaact ttctcaccac attggcctct gtgttcagca 360
ccctgcgcac cgccacncct ttgtgctggg a 391

<210> 433
<211> 420
<212> DNA
<213> Homo sapien

<220>
<221> misc_feature
<222> (1)...(420)
<223> n = A,T,C or G

<400> 433
ctgtagcttc tgtgggactt ccactgctca ggcgtcaggc tcagatagct gctggctgcg 60
tacttgttgt tgctttgttt ggaggggtgt gtggtctcca ctccgcctt gacggggctg 120
ctatctgcct tccaggccac tgtcacggct cccgggtaga agtcacttat gagacacacc 180
agtgtggcct tgttggcttg aagctcctca gaggagggcg ggaacagagt gaccgagggg 240
gcagccttg gctgacgtag gacggttagt ttggnccctc cgccgaatgc cgcanttcta 300
ctgtcccaca cctgacagta atagtcancc tcattctcgg cttgggctct gctgatggtc 360
aggggtggccc gtgntccccg agttggagcc agggaatcnc tcagggatcc canagggccn 420

<210> 434
<211> 239
<212> DNA
<213> Homo sapien

<220>
<221> misc_feature
<222> (1)...(239)
<223> n = A,T,C or G

<400> 434
ccaaccanga gagaagggat cgcttggtgc ccagggccca ccaggagctc caggcccact 60
tgggattgct gggatcactg gagcacgggg tcttgtagga ccaccaggca tgccagggtcc 120

taggggaagc cctggccctc aggggtgtcaa ggggtgaaagt gggaaaccag gagctaacgg 180
tctcagtggga gaacgtggnc cccctggacc ccagggtctt cctgggtctgg ctggtnacg 239

<210> 435
<211> 415
<212> DNA
<213> Homo sapien

<220>
<221> misc_feature
<222> (1)...(415)
<223> n = A,T,C or G

<400> 435
ctgtccaatg gcaacaggac cctcactcta ttcaatgtca caagaaatga cgcaagagcc 60
tatgtatgtg gaatccanaa ctcagtgagt gcaaaccgca gtgaccagc caccctggat 120
gtcctctatg ggccggacac ccccatcatt tcccccccag actcgtctta cctttcggga 180
gcaaacctca acctctctg ccaactcggc tctaaccat cccncanta ttcttggcgt 240
atcaatggga taccgcagca acacacacaa gttctnttta tcgccaaaat cagcccaat 300
aataacggga cctatgcctg tttagggntn taacttggnt actggccgca anaattccat 360
agtcaagagc atcacagnct ctgcatntgg aacttctcct ggctntcaga cctgn 415

<210> 436
<211> 152
<212> DNA
<213> Homo sapien

<400> 436
ccaggattga caggccatcc attcacagcc aggagatgct gggccagtc ctccaagagg 60
tctccgtcat ggcagtgatg aaaacctaac aggggtggccc cctgtgccag ctcaggtgac 120
tggagcccga gggcctgaca ggttcccagc ag 152

<210> 437
<211> 174
<212> DNA
<213> Homo sapien

<400> 437
ccagggtactg gcacatcatg ctctggatgg ggggtgggtg gtccctgtaag cagagaaaca 60
ggaaattgtc gtagtcagta tcgagcagct gtggcctcgt tcgccaccgt atagttgatc 120
ttgaacttct ttggattctc agtcttctct ccaaggacct tcttctcaac acag 174

<210> 438
<211> 485
<212> DNA
<213> Homo sapien

<220>
<221> misc_feature
<222> (1)...(485)
<223> n = A,T,C or G

<400> 438
ccacggccct ctcggccctc tcgctgggag cggagcagcg aacagaatcc atcattcacc 60

```
<210> 439
<211> 317
<212> DNA
<213> Homo sapien
```

<400> 439						
gtctt	cccctccatc	gtggggcgcc	ccaggcacca	gggcagtgat	ggtgggcatg	60
gaagg	attcctatgt	gggcgacgag	gccagagca	agagaggcat	cctcacctg	120
cccca	tcgagcacgg	catcgncacc	aactgggacg	acatggagaa	aatctggcac	180
cttct	acaatgagct	gcgtgtggct	cccgaggagc	accccggtgct	gctgaccgag	240
cctga	accccaaggc	caaccgcnag	aagatgacc	agatcatgtt	tgagaccttc	300
cccgag	ccatgta					317

<400> 440						
aagac	ttcccagggga	agatgcttgg	ctctctgctc	caaggtgggc	catggtatag	60
tcgaa	gggcttgtgg	ctggggtgat	cccagggggc	attgctcaaa	gtgcacagga	120
cagca	gggtcaggcg	agttcctgtt	ccaggggacat	caggagggag	ggtagaagcc	180
agtgt	gcgaggctgc	tgggatgagg	gagctcaggg	gctaccagct	aaccagcctc	240
aatgg	tttctccatc	cttgggtctg	tagtcagcaa	taccttgcaa	cagtgggggtg	300
gtctc	ggagaagctg	ccagaactcc	ctttctcc			338

```
<210> 441
<211> 505
<212> DNA
<213> Homo sapien

<220>
<221> misc_feature
<222> (1)...(505)
<223> n = A,T,C or G
```

<400> 441

ccacacagan	tcaccaagcc	acagacttgt	cttccacaag	cacgttctta	tcttagccac	60
gaagtgacca	agccacacgt	actaaagggt	gaactcaaag	atatgtacag	ggtattaaac	120
aaataccaag	gggaacagtt	aacttcaata	caaggtcgaa	atcagcaaca	agttctacaa	180
tccagnctg	atatcagata	caagcttcaa	ggacaatttc	ttttcgaagg	cttattccag	240
tttcgngagg	ctagcatgag	gtgtgtgcat	ttgccagggg	caaatttcta	ttctcaatta	300
acccatgcag	caaatgctac	ncatggtgcn	gagtcggttt	agaagcattt	gcggtggacg	360
atggaggggc	ccgactcgtc	ttactcctgc	ttgctaatec	acnngngctg	gaaggnggac	420
agtgaggcca	cggatggagc	caccnatcca	cacogagtnc	ttgcgctctg	ggggtgcgat	480
natnttgatc	ttcatggtgc	tgggc				505

<210> 442

<211> 386

<212> DNA

<213> Homo sapien

<220>

<221> misc_feature

<222> (1)...(386)

<223> n = A,T,C or G

<400> 442

cgccaggtga	tacctccgcc	ggtgacccag	gggctctgcg	acacaaggag	tctgcatgtc	60
taagtgctag	acatgctcag	ctttgtggat	acgcggactt	tggtgctgct	tgacgtaacc	120
ttatgcctag	caacatgccca	atctttacaa	gaggaaaccg	taagaaaggg	cccagccgga	180
gatagaggac	cacgtggaga	aaggggtcca	ccaggccccc	caggcagaga	tggtgaagat	240
ggtcccacag	gccctcctgg	tccacctggt	cctcctggcc	cccctggtct	cgatgggaac	300
tttgctgctc	agtatgatgg	aaaaggaggg	nggacttggc	cctggaccaa	tgggcttaat	360
gggacctana	ggcccacctg	gtgcag				386

<210> 443

<211> 404

<212> DNA

<213> Homo sapien

<220>

<221> misc_feature

<222> (1)...(404)

<223> n = A,T,C or G

<400> 443

cctccctctc	agagcttgcc	ccagggactc	tctggccctc	agggttcaat	gtattctgac	60
caaggccaag	ctttcctggg	gctcagggaa	aatcacactt	tgctaccoga	agctgtatoc	120
cctcagatgc	caggaaggcc	gtgatcatct	gactccaccc	tctgagaca	cattctctoc	180
ctgactgtcc	tggtctaagt	cagcggagca	ccttaggatg	gaggggtgga	ggcgaggcca	240
ngatgcagcc	tctgtgaaca	ggtgcctgga	ggctgggaaa	tgaccctgag	agggcaggac	300
acagcnaccg	ngggcttaag	gtgagggngg	agagcaagnt	tggcccaactt	tacaattcta	360
gntcagagcc	anccctaac	atggngggca	tttattcatt	tcgg		404

<210> 444

<211> 318

<212> DNA

<213> Homo sapien

<220>
 <221> misc_feature
 <222> (1)...(318)
 <223> n = A,T,C or G

<400> 444
 catgggctat agtgcgctat gttgatctgg tgttcatgct aagttccgca tcaatatngc 60
 gacttcttng gagtggggga ccaccangtt gcctaaggag ggggtgaacct gcctacgttg 120
 gaaatagagc tgggtcaaac tcctgtgctc atcagtagta gaattgcacc tgtgaatagc 180
 caccgccctc cagcntgggc aacatagcaa gacctgcct cttaagataa aaattggaaa 240
 aacttggtan gaaaaaaagg ctgtttgggtc taaanaagtc tggatngggg ataatgaca 300
 cnaancatc atgactnt 318

<210> 445
 <211> 418
 <212> DNA
 <213> Homo sapien

<220>
 <221> misc_feature
 <222> (1)...(418)
 <223> n = A,T,C or G

<400> 445
 ccagtccaac ctgctcctca ttattgtata aatgagcaga atcaatatgg cggaagccag 60
 cttcaattgc caatttgggt gcctctaaag ctttactttt aggaacctct gcaggcgcat 120
 aggtgccaaa tcccaggaca ggcataagt gaccatcatt cagcttcaca cactgatatt 180
 tcgaatccat ttctgtcact agcctggctg gcaaatgttt ctttcttctc ccctcacagg 240
 ctataagagc aatgagctgg caacgcccct gagcacactg tctgctgntt aaccaatggc 300
 atgtgagagg agggacagag gcagtcttac acaagctgtg ataaaaattg catncagttc 360
 aaccagtttc ttacnttatt ctaatngna ggaagtgtgn gaagagcaca aagtcaga 418

<210> 446
 <211> 361
 <212> DNA
 <213> Homo sapien

<220>
 <221> misc_feature
 <222> (1)...(361)
 <223> n = A,T,C or G

<400> 446
 ctgtccaatn acaacaggac cctcactcta ctgagtgtca caaggaatga tgtaggaccc 60
 tatgagtgtg gaatccanaa cgaattaant gttgaccaca gcgacccagt catcctgaat 120
 gtcctctatg gccagacga cccaccntt tccccctcat acacctatta ccgtccaggg 180
 gtgaacctca gcntctctg ncatgcagcc tctaaccacac ctgcacagta tccttggtg 240
 attgatggga acntccagna acacnacaca agagctcttt atctccancn tnactganaa 300
 gaacagcgcg actctatncc ttccaggggg ggggggtggg gnntgnggac cttncggggc 360
 c 361

<210> 447
 <211> 321

<212> DNA
<213> Homo sapien

<220>
<221> misc_feature
<222> (1)...(321)
<223> n = A,T,C or G

<400> 447
ccagganant ggttccccaa aggggacctc acccgccccg agctctggag ccgctgacgc 60
tcgcatccag gacatttgag atgggaatcc aaataggcta cttgnaaaaag acgtgctgca 120
ngcagccctg gagagactca tggagttcat tgtacattac tccatctacc gaggcagcgc 180
atggcatgac tnaacggctt gnaacaaaaca canaaattac caccacaaac attcaggaac 240
caaataataat ctgctatggt cacaccacag acaatgcagg aagaggcttt ttattgctng 300
ngtgngtttt caaatcatgt t 321

<210> 448
<211> 325
<212> DNA
<213> Homo sapien

<220>
<221> misc_feature
<222> (1)...(325)
<223> n = A,T,C or G

<400> 448
ccagcttcaa ctttttagta tagaagatac aggatcacaa aaaggagact acgctttgca 60
aacatagcat caaaattcaa cttttctctt tgcagtttat ccatggngtc agcatacett 120
gcaagggaag ctacttacat caaataactt ttctatatac atttcctcat tgaccttttc 180
tcaaagaata tcttggtttt gccgaacaaa cataatatag gngtctgcc gatccattcc 240
tggtttctgt ngtgaaggaa aagcaggggg aacaaaataa tatcagggtc tcaatngtga 300
nattattatt taatcatacc ctgan 325

<210> 449
<211> 123
<212> DNA
<213> Homo sapien

<220>
<221> misc_feature
<222> (1)...(123)
<223> n = A,T,C or G

<400> 449
cattaatntt ggaagcgatg gtgtggatta catcagtgtt agggcatggt gtggatatta 60
ttacattann attggaagcg atgggtgtgga ttacatcagt gatagggcac ggtgtggata 120
tta 123

<210> 450
<211> 328
<212> DNA
<213> Homo sapien

<220>
 <221> misc_feature
 <222> (1)...(328)
 <223> n = A,T,C or G

<400> 450
 ctggcaattt tgagctgccg gttatacacc aaaatgttct gttcagtacc tagctctgct 60
 cttttatatt gctttaaatt tttaaagaaa ttatatgtca tggatgtggt tatttgtgca 120
 tattttttta caatgcccaa tctgtatgaa taatgtaaac ttcgattttt ttttaaaaaa 180
 attagatttt agctggagct tttgactaat gttaaagtaaa tgccaaacta ccgacttgat 240
 ngggatgttt ttgtaangtt aatttttctaa gactttttca catccaaagt gatgctttgc 300
 tttgggtttt aactgtttca acntnggn 328

<210> 451
 <211> 209
 <212> DNA
 <213> Homo sapien

<400> 451
 ctgccttggt tcaacagaca tgcaaagatc ctaggagaca gtcccatag accttcagac 60
 attaaaaagg gagccgtaca gtttgtttga agcacttcgt cttaccatt tatgcagggg 120
 ccccaggaaa cttacacaca gccagaatga gggtcccaaa ggacttacat taattatggc 180
 tcttgcttcc tttcacaaat gagctgagg 209

<210> 452
 <211> 457
 <212> DNA
 <213> Homo sapien

<220>
 <221> misc_feature
 <222> (1)...(457)
 <223> n = A,T,C or G

<400> 452
 ctgtctantc ctttcaagag ctgtttatag aagcttgaga atggggtaaa aatttctgct 60
 agcaaaatca agttcttttt gaaattttat cagtaatcca gaatttagta gtocatgcct 120
 tctcactcag catttagaaa taaaaatgtg gtttcttaaa cgtatatcct ttcattgata 180
 tttccacatt tttgtgcttg gatataagat gtatttcttg tagtgaagtt gttttgtaat 240
 ctactttgta tacattctaa ttatattatt tttctatgta ttttaaagn atatggctgt 300
 ttaattcttg aagcattttg ggcttaagat tgccagcacc acacatcaga tgcagtcatt 360
 gttgctatca gtgtggaatc tgatagagtc tngactccgg ccacttggag ttgtgnactc 420
 caaagctaag gacagtgatg aggaagatgg catgtgg 457

<210> 453
 <211> 277
 <212> DNA
 <213> Homo sapien

<400> 453
 ccaattgatt tgatggtaag ggagggatcg ttgacctcgt ctgttatgta aaggatgcgt 60
 agggatggga gggcgatgag gactaggatg atggcgggca ggatagttca gacggtttct 120
 atttcttgag cgtctgagat gttagtatta gttagttttg ttgtgagtgt taggaaaagg 180
 gcatacagga ctaggaagca gataaggaaa atgactacga gggcgtgatc atgaaagggtg 240

ataagctctt ctatgatagg ggaagtagcg tcttgta

277

<210> 454
<211> 198
<212> DNA
<213> Homo sapien

<400> 454
gttaaaagat agtaggggga tgatgctaata aatcaggctg tgggtgggtg tgttgattca 60
aattatgtgt tttttggaga gtcattgcag tggtagtaata ataattgttg ggacgattag 120
tttttagcatt ggagtagggt taggttatgt acgtagtcta ggccatatgt gttggagatt 180
gagactagta gggctagg 198

<210> 455
<211> 608
<212> DNA
<213> Homo sapien

<220>
<221> misc_feature
<222> (1)...(608)
<223> n = A,T,C or G

<400> 455
ctgagcaagc taaggaccag gggcaactag accctaataa tnggtacttt tgaaaatgat 60
acaaactacc ttggttgtaa gaagtgcagg ttgaacactt taggagaaca gtcttcaaac 120
tggcaattca aaatttccca ttatatgtga ataaaattgg aaggatgtta aatgtccatg 180
gaaagttact cttgttaagt aggatgcctt atactgaggc tttanaatga aagtacactt 240
cacaaatgga atagtgaaca taaattacca gaagtcaaga taatagtcac actagtaagg 300
taagcaaggc aaattccctt atacacaaaa attattttga tgaccttttt caataatgaa 360
tctgaaatga agtggtttta aaagctccct aaacacaaaa cgaacataaa actgcttaat 420
aacttttagag ctcatgtaat attcttgctg aaacacagta ctgaaattac cagcgaaatg 480
atggaatatc tttaaagcag gncactcngt ataactctgga ataatttcac ttgctaactt 540
ttaagaagta ttctctggac tataaatcnt gggcaaatag acttccactt tattattacc 600
ccaaatta 608

<210> 456
<211> 467
<212> DNA
<213> Homo sapien

<220>
<221> misc_feature
<222> (1)...(467)
<223> n = A,T,C or G

<400> 456
cctggacctg tgtaaacctt caaacactct tttttacatt aggtcgtgaa gttaaatttt 60
ttactgtttc tgtgctacag actcttcaaa gggaaatagt taagtcaatt tcaaagaaaa 120
tgaccagcac attttttaaaa cattagaaat gatttgactt tgactatcta ctgccccaaa 180
aagggttaagg aatttgtaat gagaagctaa aaactttaag gaattttaag gaactcaaaa 240
caaaaactca ttaaagttaa ttaaagttaa ttctacaaat aaagcctctt aatacatctt 300
tataatagtc acttaagact taaattcaaa cactagcaaa ccacaaaatc agactgtntg 360
actgacatcc aaaagataaa tataaatcaa aatccgaccc cagcattagc caaggggtag 420

467

<400> 457

```
<210> 458
<211> 445
<212> DNA
<213> Homo sapien
```

```
<220>  
<221> misc_feature  
<222> (1)...(445)  
<223> n = A,T,C or G
```

<400> 458

gaaaaatata	aagccaaaaa	ttggataaaa	tagcactgaa	aaaatgagga	aattattggt	60
aaccaatttta	ttttaaaagc	ccatcaattt	aatttctggt	ggtgcagaag	ttagaaggta	120
aagcttgaga	agatgagggt	gtttacgtag	accagaacca	atttagaaga	atacttgaag	180
ctagaagggg	aagttggtta	aaaatcacat	caaaaagcta	ctaaaaggac	tggtgtaatt	240
taaaaaaaaac	taaggcagaa	ggttttttgga	agagttagaa	gaatttggaa	ggccttaa	300
atagtagctt	agtttgaaaa	atgngaagga	ctttcgtaac	ggaagtaatt	caagatcaag	360
agtaattacc	ancttaatgt	ttttggcntt	ggactntgag	ttaagattat	tttttaaattc	420
ctgaggacta	ncattaatgg	gacag				445

```
<210> 459
<211> 426
<212> DNA
<213> Homo sapien
```

```
<220>
<221> misc_feature
<222> (1)...(426)
<223> n = A,T,C or G
```

<400> 459

cctatgatan	cttctctagc	tatcatactc	caatcagcaa	aaaatgagaa	aatgttgaga	60
aatagaagat	aattcctcat	ttaaggccac	cttctagaat	ttgtgcttaa	gattctgctt	120
tcttctcatg	ggccagcact	tcggcaactg	gcaaaaatta	ggtgtacagg	gatctaggta	180
atactgttta	tttgagcaat	aatatattgt	gctaacgttc	aggcatccta	ttactgagaa	240
ataaggqaaa	atgagtgtaa	agtacaacta	agagtctcgg	cgacagggaa	aaataccatc	300

agttaaatat ccatagtcct agagcattta tgtaaaactg caatntgaat cctgcaatac 360
 atnttggctt tttccctcag tgataccatg tgaggaagn ngctctgtca aggcgggccc 420
 gataga 426

<210> 460
 <211> 348
 <212> DNA
 <213> Homo sapien

<220>
 <221> misc_feature
 <222> (1)...(348)
 <223> n = A,T,C or G

<400> 460
 ccaaatttta aaatgttatt tttcatatca tttataacct tgtcacaatc cacttaaaga 60
 agtttgggta tatttcactg aaaattttct tccagagtag gttttttttc gtgggttggg 120
 gggtaacttt actacaatta gtaagtntgg tgcagaatth catgcaaag aggagtgag 180
 cagngtgata atttaaacat atntaaacaa aaacaaaaaa aatgaatgca caaacttgct 240
 gctgcttaga tcaactgcagc ttctaggacc cggtttcttt tactgatnta aancaaaaac 300
 aaaaaaanta annacnttgt gcctgaaatg aancttggtt tttntna 348

<210> 461
 <211> 378
 <212> DNA
 <213> Homo sapien

<220>
 <221> misc_feature
 <222> (1)...(378)
 <223> n = A,T,C or G

<400> 461
 ccactaagac agaacggaat ctagtagaag tgcaccaatg cttcagtcct tctactcag 60
 catgggtgagc agtgggtcaat ctgtgccctg tggagtgatg ggcagataat tctggcatgt 120
 gtaaataata ataaataatt cacttggtgc aggcagtagt tctatgaatt aaaacctagt 180
 gtgtacacag tgcctacatg tgttacagcc ccacagtagg aatctacacc aaaatattta 240
 ttagaaggaa tttgggtccgt actacatcac gctttccgga gggtaaaaaa taaagtccat 300
 ctatagacat ttcaccacag acccagagac tgagtctggc taaaacctgc aaaatgtcta 360
 taacaaaagn ggatggct 378

<210> 462
 <211> 197
 <212> DNA
 <213> Homo sapien

<220>
 <221> misc_feature
 <222> (1)...(197)
 <223> n = A,T,C or G

<400> 462
 gcgaggtcca cactattaaa agctgttggg taattgaagg tgatataaaa tgactgtcnt 60
 catttggagt gngcagcaca nttacttcat gttgctcang tttanaacaa tntccctgn 120

<400> 465
ccactctttgg tagaaacctt gaaactttca ctttgcctggg ctttagcaaa gtttcctttt 60

```

acagttctgt ttatgagctt cagctactga taaagcactt cctgaacttc tctattatca 120
tagngaccct ctgaataacc tgagtgactg gctcggcaat tcgctttata accattctta 180
ttcccaaagt tggagcacat aaacatcttag atgtcttttc ctgtaaaata ttctagacat 240
ttaccctaac tctagttcaa catatactca acttgcaactg tatactctccc tgcttttttg 300
agacagagaa gaaattcagg aggtgnoccc tctccagagt ttctctgttg gaaagcagcn 360
atcaagaanc ctttaaaaaa ttggtgtnaa gctntgcnc ctgcagaaat gcntngcccc 420
acattattct tctggggnaa agna 444

```

<210> 466

<211> 381

<212> DNA

<213> Homo sapien

<220>

<221> misc_feature

<222> (1)...(381)

<223> n = A,T,C or G

<400> 466

```

cctactatgg gtgttaattt tttactctct ctacaagggt ttttcctagt gtccaaagag 60
ctgttcctct ttggactaac agttaaat tacaaggggat ttagagggt ctgtgggcaa 120
atttaaagtt gaactaagat tctatcttgg acaaccagct atcaccaggt tcggtaggtt 180
tgctgctctt acctataaat cttcccacta ttttgctaca tagacgggtg tgctctttta 240
gctgttctta ggtagctcgt ctggnttcgg gggctcttagc tttggctctc cttgcaaagt 300
tatttctagt taattcatta tgcannaggt ataggggnta gtccttgcta tattatgctt 360
ggttataatt tttcatcttt c 381

```

<210> 467

<211> 95

<212> DNA

<213> Homo sapien

<220>

<221> misc_feature

<222> (1)...(95)

<223> n = A,T,C or G

<400> 467

```

cctatanatt ntggnttgta tactgggtcc tgaaaaccct cttggngctc tgtttttaag 60
gagctgaanc caanganccg caataataat acttt 95

```

<210> 468

<211> 224

<212> DNA

<213> Homo sapien

<400> 468

```

cagtgggtct ctgatgcctt gcctgcagca gaaggaggga gcagagatca agaggaagga 60
aaaaatcata tgtacttatt tgaaggtaaa gattattcta aagagcccag taaggaagac 120
agaaaatcat ttgaacaact ggtaaacctt cagaaaaccc ttttggagaa agctagtcaa 180
gagggccgat cactccgaaa taaaggcagt gttctcatcc cagg 224

```

<210> 469

<211> 416

<212> DNA
<213> Homo sapien

<400> 469
ctgagttcta gttcaaaagc tttatcctta acttcgtcat gtactatgta aattctagaa 60
tagaaaaggg aaaggtaaga ttttggtaac ctccaaacat tgaagtagtt cacagaccca 120
aagtcagtac aaattagaat gtccatccat aataaaagta tctataaaat tacacagaca 180
cattctacat agtatTTAAC attagagaag acaaattaca cagggactga aataaaatga 240
aacatctact ctcccgacaa atgttgaata tacctaatac acccaagttc agtttatttt 300
tgcacattgc ttttagagata taacttggct gggcacagtg gtcacacct gtaatcccaa 360
cactttggga gaccaaggcg gatggatcac ttgaggtcag ttcgagacta gcctgg 416

<210> 470
<211> 376
<212> DNA
<213> Homo sapien

<400> 470
caccttttaa ctgtatcaca aagtctgttg ctgtgggttac agcctttgtt tccagtgatg 60
ttttgtccat gctttccccc aacccttaac aatgggttact caaaagaatg aaataatgag 120
tcattcattc gggaatatgt taaaaatccc ctctttatca ttacatttca ctgcttagaa 180
actaggctgt aattcaaggc aacagttaag tctgagaact gttaaaaaaa tctttgattt 240
tttttcattt ttaagaaaaa cctgcctatt taattgttca gacttgtaag aggttcttca 300
attacatcct ttttggttaa tgtattattt ctggaacaag tagataaaat tctacgcagt 360
aagcataata aaatc 376

<210> 471
<211> 357
<212> DNA
<213> Homo sapien

<400> 471
ggcttcgtat aatgggttctt ttgtcacccc tgatcgacga tttcgctacc cgtacaactc 60
tgacaaggga acgaaatgct tctgtgtatt cacctagtgg tctgtgaac agaagaacaa 120
caactccacc ggatagtggg gtactgtttg aagggttagg catttcaaca agacctagag 180
atgttgaaat tcctcagttt atgagacaga ttgcagtaag gaggccaact acggcagatg 240
aaagatcttt gcggaaaatt caagaacaag atattattaa ttttagacga actctttacc 300
gtgctggtgc tcgagttaga aatattgaag atggtggccg ctacagggat atttcag 357

<210> 472
<211> 557
<212> DNA
<213> Homo sapien

<220>
<221> misc_feature
<222> (1) ... (557)
<223> n = A,T,C or G

<400> 472
cngagatgac atttacaatc tcttgaaang cagcagatgg cactctggtg cttcctatga 60
agcaacatgc ttgaaatcaa gggccaacaa ttgtttagg aaagcaaaat atacctctaa 120
cacctacgtt taccaaaaaa gctgacatct caaactctga gttgttgaga ctcaaatttc 180
tcatcccaa agaagcctat tacggtagtg tgntggatgc tttttgtatc tctgataggc 240

```

aggcactata atgggggggaa atacttctga ataaaaacat tggctgtctt gcaactgtgc 300
atataatgtc tattcaaggg ggcagtgtgc ctagcatgat cctgaaatgt tgagataaaa 360
ggaagtgtgc attaaagcac tatttgtctt atatgaaaag agtgactcta tcttccagta 420
aacaagantt cctgcaatga aaaagaaatt ttttccttca ttatctataa actatacaaa 480
ataaccttcc tttttaacct aagactcaaa cattnatatt tgattttatt ctatttgata 540
ccaattggta tgtccag 557

```

```

<210> 473
<211> 264
<212> DNA
<213> Homo sapien

```

```

<400> 473
cctccatcaa cagaaaggat aaagacccct tcgggtctcc tcattaattc tgaactggaa 60
aagccccaga aagtccggaa agacaaggaa ggaacacctc cacttacaaa agaagataag 120
acagttgtca gacaaagccc tcgaaggatt aagccagtta ggattattcc ttcttcaaaa 180
aggacagatg caaccattgc taagcaactc ttacagaggg caaaaaaggg ggctcaaaag 240
aaaattgaaa aagaagcagc tcag 264

```

```

<210> 474
<211> 165
<212> DNA
<213> Homo sapien

```

```

<400> 474
aattcagctt ccagaggccc ttattagtcc ttgttgacag aaacatagat ttggcaactc 60
ctttacatca tacttggaca tatcaagcat tgggtgcacga tgtactggat ttccatttaa 120
acagggttaa tttggaagaa tcttcaggag tggaaaactc tccag 165

```

```

<210> 475
<211> 417
<212> DNA
<213> Homo sapien

```

```

<220>
<221> misc_feature
<222> (1)...(417)
<223> n = A,T,C or G

```

```

<400> 475
aagttctctt cttgttttaa acacattcct gataacttct aaagatgacc aaaataaaac 60
agaatatcta cagagatcat tttctgaatt ttttgtacat ccaaggataa caacataaaa 120
aaaataaaac tggacagcat tccacatcca agtgcacaga accatttttg caagattaaa 180
taatgtaaac attgggaaca gccaaatcag cgaagaatgc caacacctca aaacacctgg 240
tgttgccgct tcattaagtg gttcaaaaac cagatctata attgcgcaat attcacgta 300
tataaaaaga aatggatatt aattttgaca aatagctgca actgagactt ctttttattt 360
ctttatatgn gnatatagtg aatttttatt attttttaaa ttttatttat tttttta 417

```

```

<210> 476
<211> 321
<212> DNA
<213> Homo sapien

```

```

<220>

```

<221> misc_feature
 <222> (1)...(321)
 <223> n = A,T,C or G

<400> 476
 catttaataa caaaaacaac ctgtacggaa aaccnaagg caaccacata gcatatgtaa 60
 aatgtgcaaa tacacttttaa aatgcangtt attctatagc anttgcaaga tagaatttca 120
 ctgtaattag ggaatctagc tcacctaac ttaatagnct tttgcatgtn tagacaatgc 180
 aattctacaa ggnacnactc agcgttgatg cttaaagtatg aaacacatcc tcagattatt 240
 catccgaaaa tattaaaata gcntcatggt ttattattct ttaatgagtc ntgagctcat 300
 ttctaaagct tcataaagca t 321

<210> 477
 <211> 546
 <212> DNA
 <213> Homo sapien

<220>
 <221> misc_feature
 <222> (1)...(546)
 <223> n = A,T,C or G

<400> 477
 gctgtggtta tattgtaaat gaagcatcta acatgtgcac aacttgcaac aaaaactcct 60
 tggactttta atctgtcttt ctacagttcc atgtgtgat tgatctgact gatcacacag 120
 gcacccttca ttctgtagt ctacaggaa gtgttgctga ggagactttg ggctgcacgg 180
 tacatgagtt tcttgcaatg acaaatgaac agaaaacagc attaaagtgg caattcctct 240
 tggaaagaag caaaatttat ttaaaattcg ttctatcaca cagagcaagg agtggattga 300
 aaattagtgt actctcgtgc aagcttgagc atcctactga ggcaagcaga aacttgtctg 360
 gacaaagaca tgtttaaaac ggtctatcat tttgaactct ggaaaagtat aagagtttta 420
 actcccttta aaatggaata ttaatttgaa aattatgggg aaaattgcat tttgtttaca 480
 tgtggtgaac atgtttctag aaattggtat ggcgggaagg gggctgggtg agtctgaagg 540
 acctcn 546

<210> 478
 <211> 100
 <212> DNA
 <213> Homo sapien

<400> 478
 aagaaaagtg gtaaaatcaa gtcttcttac aagagggagt gtataaacct tggttgtgat 60
 gttgactttg attttgctgg acctgcaatc catggttcag 100

<210> 479
 <211> 508
 <212> DNA
 <213> Homo sapien

<220>
 <221> misc_feature
 <222> (1)...(508)
 <223> n = A,T,C or G

<400> 479

```

gnnttccaaa ttcttctaac tcttccaaaa gccttctgcc ttagtttttt tttaaattaca    60
ccagtccttt tagtagcttt ttgatgtgat ttttaaccaa cttccccttc tagcttcaag    120
tattcttcta aattggtoct ggtctacgta aacaccctca tcttctcaag ctttaccttc    180
taacttctgc accaccagaa attaaattga tgggctttta aaataaattg gttaccaata    240
atttcctcat tttttcagtg ctattttatc caatttttgg ctttatattt ttctatcttc    300
tatacttctc caatacttgt cttagcttgt ttttcatttt ctatctgaaa ctcttgacaa    360
tatcttctaa tttccctatc ttctctatct ttttcttcgc cttcccgtac ttctgcttcc    420
agntttccac ttcaaacttc tatcttctcc aaattgttca tctaccact cccaataatc    480
tttccatttt cgtgtagcac ctggncag                                     508

```

```

<210> 480
<211> 81
<212> DNA
<213> Homo sapien

```

```

<400> 480
ggtgcccttt tcctaacact cacaacaaaa ctaactaata ctaacatctc agacgctcag    60
gaaatagata aggaaaatga c                                           81

```

```

<210> 481
<211> 306
<212> DNA
<213> Homo sapien

```

```

<220>
<221> misc_feature
<222> (1)...(306)
<223> n = A,T,C or G

```

```

<400> 481
tcgccttcgg ccgccgggca ggtaggggn acaagacgct acttccccta tcatagaaga    60
gcttatcacc tttcatgac acgccctcat agtcattttc cttatctgct tcttagtcct    120
gtatgccctt ttctaacac tcacaacaaa actaactaat actaacatct cagacgctca    180
gggaatagaa accgtctgaa ctatcctgcc cgccatcctc ctagtcctca tcgcctccc    240
atccctacgc atcctttaca taacagacga ggtcaacgat cctccccta ccatcaaatc    300
aattgg                                           306

```

```

<210> 482
<211> 582
<212> DNA
<213> Homo sapien

```

```

<220>
<221> misc_feature
<222> (1)...(582)
<223> n = A,T,C or G

```

```

<400> 482
ggggggaaca gtcattatac attatttaga ctcatctcct cttccagtgc ccttatgatt    60
atttcctacc tttaccattg atcttaaaact gngcaggcta aaaagaggaa ccagaactcc    120
cttaagcact tttaagacta tttaaaaaat aaagnnttgt tggcattgaa gagtaagctg    180
cttaagggac tgaatgaaaa gatagtaccc tttgtggctg tatgaagaga gaaactgaat    240
ttctatccaa gagaccttaa tntagcctat tagggaatta tcttcccaa aagtacaagt    300
aattttgcac tgcaggagaa ggataagtag atttgattta catcacattt tatacacacc    360

```


<210> 486
 <211> 274
 <212> DNA
 <213> Homo sapien

<400> 486
 ctgtaatat gtagttgctc cagaatgtca agggcagctt acggagatgt cactggagca 60
 gcacgctcag agacagtga ctagcatttg aatacacaag tccaagtcta ctgtgttgct 120
 aggggtgcag aaccgcgtttc tttgtatgag agagggtcaaa gggttgggtt cctgggagaa 180
 attagttttg cattaaagta ggagtagtgc atgttttctt ctgttatccc cctgattgtt 240
 ctgtaactag ttgctctcat ttttaatttca ctgg 274

<210> 487
 <211> 184
 <212> DNA
 <213> Homo sapien

<220>
 <221> misc_feature
 <222> (1)...(184)
 <223> n = A,T,C or G

<400> 487
 tggcaccaag attctcagct cacggtacca gcatctgatt gtcggactac ctgctgcttt 60
 ccctgatatt tatacatgat attcgnaaaa tgtaaagaag ctattattca tacagacatc 120
 tagagaagga gngaagnttt taaaaaaata aaaaaatact tatttcaagc tttagctgtg 180
 ttct 184

<210> 488
 <211> 393
 <212> DNA
 <213> Homo sapien

<400> 488
 ctgcattttt attgcatct gcagatgaac tggaaaatct cattttacaa cagaactggg 60
 acagacgacc accatattca ctgaggtcta aatttgcagt ttccactaat gacattttga 120
 tttcccaaca gagatacttc tggctcttact gcacagtctt ttaagagaaa tacttccatt 180
 atgccacatt gtccttgatc cgtaagtgat gtgttaaggt gcttcaaagg aactctgacc 240
 tctgaagtac ttgagctact ttagtatgtc cagcctattg ctttttggtt tagtgtgtca 300
 ccataaatat caggggcata aaaggctatc tattcttaat tcaaggataa aacagaagaa 360
 gcttggtgga taaaacaata gttcaagatc cag 393

<210> 489
 <211> 607
 <212> DNA
 <213> Homo sapien

<220>
 <221> misc_feature
 <222> (1)...(607)
 <223> n = A,T,C or G

<400> 489
 gtgcttatgt acttaagggg aactactcta actgggtgaa gagtangatg aagcatccat 60

<400> 492
ctccacctta ctaccagaca gccttagcca aaccatttnc ccaaataaag tataggcgat 60
agaaattgaa acctggcgca atagatatag taccgcaagg gaaagatgaa aaattataac 120
caagcataat atagcaagga ctaaccccta taccttctgc ataatgaatt aactagaaat 180
aactttgcaa ggggagccaa agctaagacc cccgaaacca gacgagctac ctaagaacag 240
ctaaaagagc acaccgtct atgtagcaaa atagtgggaa gatttatagg tagaggcgac 300
aaacctaccg agcctggtga tagctggttg tccaagatag aatcttagtt caactttaaa 360
tttgcacaca gaacctcta aatccctctt taaatttaac tgttagtcca aagaggaaca 420
gctctttgga cactaggaaa aaaccttgta gagagagtaa aaaatttaac acccatagta 480
gg 482

<210> 493
<211> 207
<212> DNA
<213> Homo sapien

<220>
<221> misc_feature
<222> (1)...(207)
<223> n = A,T,C or G

<400> 493
cataaatatt atactagcat ttaccatctc acttngngga atgctagtat atcgctcaca 60
cctcatatcc tccctactat gcctagaagg aataatacta tcaactgttca ttatagctac 120
tctcataacc ctcaacaccc actccctctt agccaatatt gtgcctattg ccatactagt 180
ctttgccgcc tgccaagcag cggtagg 207

<210> 494
<211> 283
<212> DNA
<213> Homo sapien

<220>
<221> misc_feature
<222> (1)...(283)
<223> n = A,T,C or G

<400> 494
ccaattgatt tgatggtaag ggagggatcg ttgacctngt ctgttatgta aaggatgcgt 60
agggatggga gggcgatgag gactaggatg atggcgggca ggatagttca gacggtttct 120
atttcttgag cgtctgagat gttagtatta gttagttttg ttgtgagtgt taggaaaagg 180
gcatacagga ctaggaagca gataaggaaa atgactatga gggcgtgatc atgaaagggtg 240
ataagctctt ctatgatagg ggaagtagcg tctttagtagac cta 283

<210> 495
<211> 590
<212> DNA
<213> Homo sapien

<220>
<221> misc_feature
<222> (1)...(590)
<223> n = A,T,C or G

<400> 495
 tatgtatatata atttttcttag ttactagcat agagaaatta ctgattttaa aaaacatttc 60
 aaattctagc atgttgtagg attctattgc cctttctaaa aagtacatct tgcttatccg 120
 atttctaaca aaactattta atttgaagaa gggagaatga atttggataa aaagcaaaaa 180
 tttaaaggta ctcaaattta ggcaaaccat taaagcaatc ttagtttaca gttaattggg 240
 tagaatgggc aacactttct tcagggttagt tcatggagtg gatatgcatt gatagaacaa 300
 cttagagatg cttttacagt tgagaaagct cattatatct gttatcttta agaatcagct 360
 tatttatttc atatgtttgt tctttaagaa gaccaaagag ccctgcaaat gaatgttgat 420
 ttgttttttt gtttgtttaa tatttttgta gagataagat ctcactttgt tatgttgccc 480
 aggctggtct caaactctca acttgaagtg atctgccac ctcagcctcc caaagtgggtg 540
 ggattacagg catgagccac cgcacctgga cctgcccggg cggncgctcg 590

<210> 496
 <211> 307
 <212> DNA
 <213> Homo sapien

<220>
 <221> misc_feature
 <222> (1)...(307)
 <223> n = A,T,C or G

<400> 496
 ggagattagt atagagaggn anacnttttt tcngnatatt tggtcacatg gataagtggc 60
 gctggcttgc catgattgtg aggggtagga gccaggtagt tagtattagg aggggggnng 120
 ttagggggtc tgaggagaag gttggggaac agctnaatag gttgttngnt gatttggnnta 180
 aaaaacanta gggggatgat nctaataatt antgctgtgg gtggttgtgn tgattcaaata 240
 tatngtcttt ttcggagann catgtcangt ggtagtaaata ataattgttg ggaccattan 300
 ttcttan 307

<210> 497
 <211> 216
 <212> DNA
 <213> Homo sapien

<220>
 <221> misc_feature
 <222> (1)...(216)
 <223> n = A,T,C or G

<400> 497
 cattttctctc ttgggtttctt cagttaagtc aaannngnac gttctctctt ccccatatat 60
 tcatatatatt ttgctcgtaa gtgtatttct tgagctgttt tcatgttggt tatttctctgt 120
 ctngaaatg gtgttttttt ttgttggtgn tgggtttttt tttttttttt aaactnngna 180
 ccncaantt gaaaaaatgn ttntttttcc ctnaca 216

<210> 498
 <211> 375
 <212> DNA
 <213> Homo sapien

<220>
 <221> misc_feature

<223> n = A, T, C or G

gaatttcctg	gcaccttttc	tgcctagaga	agattnngtg	tgactggggt	gcctataagc	60
catatagata	caaactttta	tctctaatac	caagtcttag	agggatatat	taatagatct	120
aataaaattta	ttcttagact	tattgtttca	tgggntagtg	agtctttgct	actggagaca	180
atacagactt	gtcagttttt	ttaaaaaaa	aaaatttgcc	aagctancac	attaaanaa	240
tntcctaagg	ctntcatttt	atgaggatga	ttataaacnt	ttntgngata	aatatcacca	300
taataaaactg	ttaagtacaa	ctgcnggccc	cccttanagn	gaattcctnc	agttanaaat	360
ttatttttttt	gccaa					375

<211> 215

<213> Homo sapien

<221> misc feature

<223> n = A, T, C or G

ccacnaaaagc	agaagcttaa	agcatagtag	taaagaggnn	aaaaagaagg	acgaaaataa	60
atcatagatgc	aaggatggta	aagaagttga	cagtagtcat	gaaaaggcca	gaggtaatag	120
ttcactcatg	gaaaagaaat	taagtagaag	gttgtgcgaa	aatcggagag	gaagcttgtc	180
acaaaaaaaa	aaaaaaaaa	aaaaaaaaat	gtttt			215

<211> 489

<213> Homo sapien

<221> misc feature

 $\langle 222 \rangle \quad (1) \dots (489)$

<223> n = A, T, C or G

ccactacgat	aagcaggtag	ctgggttttg	tagtgagntt	gctccttaag	ttacaggaac	60
tctccttata	atagacactt	catttttccta	gtccatccct	catgaaaaat	gactgaccac	120
tgctgggcag	caggagggat	gatgaccaac	taattcccaa	accccgagtct	cattggtaacc	180
agcctttgggg	aaccacctac	acttgagcca	caattggttt	tgaagtgcac	ttacaaggnt	240
tgtctactttt	cagttcttta	ctttttacat	gctgacacac	acatacactg	cctaaataga	300
tctctttcag	aaacaatcct	cagataacgc	atagcaaaat	ggagatggag	acatgatttc	360
tcatgcaaca	gctttctctaa	ttatacctta	gaaatgttct	cctttttatc	atcaaattctg	420
ctcaagaag	gctttttata	gtagaataat	atcagtggat	gaaaacagct	taacatttta	480
ccatgctta						489

<211> 286

<212> DNA

<213> Homo sapien

<400> 501
 aaaaacactc aaacacagcc ttggagggag gagtcagttt taaaagactc ttataaaaagt 60
 aatatactgc tagctctgaa gaatcggagg ctaaaatcat ctcttcaagt cccaggggaa 120
 tcccaaagaa ctccagggga aggtgggatg ggccagagag ctctggaagc ttccaggtct 180
 gttgcaagcc tcacctggta cacagtaggc tcttccaggt ctgtcaggaa cccaggagcc 240
 tcccctagca cacagtaggc tcacaaaaag ggagcactgc tgctgg 286

<210> 502
 <211> 168
 <212> DNA
 <213> Homo sapien

<220>
 <221> misc_feature
 <222> (1)...(168)
 <223> n = A,T,C or G

<400> 502
 cctatgattg tgggggcaat gaatgaagcg aacagagntt cgttcatttt ggttctcaga 60
 gtttggtata attttttatt tttatgggct ttggtgaggg aggtaagtgg tagtttgtgt 120
 ttaatatatt tagttgggtg atgaggaata gtgtaaggag tatggggg 168

<210> 503
 <211> 173
 <212> DNA
 <213> Homo sapien

<220>
 <221> misc_feature
 <222> (1)...(173)
 <223> n = A,T,C or G

<400> 503
 cctttataat aaattaggca aaaggttcag tgcnnngcta tantggacaa catgaaactc 60
 cataaaaatg actggatagg gggactgctt gagacttttc ttttgggcat tactaacaga 120
 attcaaagaa attccaacca cgcttatttt tccaaattct actgaaatga gag 173

<210> 504
 <211> 310
 <212> DNA
 <213> Homo sapien

<220>
 <221> misc_feature
 <222> (1)...(310)
 <223> n = A,T,C or G

<400> 504
 tagtattcta tttaaaaatt aagttttggg gtctgtaaaa tatacaggac aatgactttt 60
 ttaaaatgta agttaatacc tcctcctcac ttgtcttaat tgaacttagg tgtttattct 120
 taaagngnga ccttgatgaa aatgttgaga tgggaagtgt tattaggcaa aacttggtat 180
 agattttctca tataactctt aattgaccct tagaatttta acaaccgcgc ctggcccaat 240
 agactgtttt ttagagtant tttaggctct cancaaaatt gaggggaaaa tacagggtgt 300
 tcccattaaa 310

<210> 505
 <211> 530
 <212> DNA
 <213> Homo sapien

<220>
 <221> misc_feature
 <222> (1)...(530)
 <223> n = A,T,C or G

<400> 505
 cctcagggaa cttacaatta tggcaaaagg ggaaggggaa gcaagcacct tcttcacaag 60
 gcatcaggag agagagagaa agagagtagg ggaaactacc ctttttaaac catcatatcc 120
 tgtgagaact ccctcagtat tagaagagca tgagggaaac cgctccata atccaatcac 180
 ctcccaccag gaccatccct caatacatgg gggttacaat tcaagatgag gttcgggtgg 240
 ggatacagat ttaaaccata tcagaatggg taatgatatt gttgtatatt accaactata 300
 atcttcttag tggtatagta caataatgta aaaaattgag taaatttggt ttctatatta 360
 ttctgttttt ggaaaacatg tatatagtca gggctgtttg tctcaagaaa atatggtaaa 420
 ctctgctgtt ttggtcactg gtgcctagaa tttggggatg tacattgggt ttgattcaca 480
 tgcacatttc cttctagtgc acagtaacta tttctaacta tttcccnata 530

<210> 506
 <211> 352
 <212> DNA
 <213> Homo sapien

<220>
 <221> misc_feature
 <222> (1)...(352)
 <223> n = A,T,C or G

<400> 506
 cttgaacgct ttcttaattg gtggctgctt ttaggcggtg ctatgggtgn taaatTTTTT 60
 actctctcta caaggttttt tcctagtgtc caaagagctg ttctcttttg gactaacagt 120
 taaatttaca aggggattta gagggttctg tgggcaaatt taaagttgaa ctaanattct 180
 atcttgga accagctatc accaggctcg gttaggtttg cgctctacc tataaatctt 240
 cccactattt tgctacatag acgggtgtgc tcttttagct gttcttaggt agctcgtctg 300
 gtttcggggg tcttagcttt ggctctcctt gcaaanntat ttctagttaa tt 352

<210> 507
 <211> 370
 <212> DNA
 <213> Homo sapien

<220>
 <221> misc_feature
 <222> (1)...(370)
 <223> n = A,T,C or G

<400> 507
 cctaactaga tcttatcaga atagggggga agggngtcgg ttcattcetta ttgagtgtta 60
 atgaccctgt aagatgtaat ttcttttatt tcattctggt acctagaaaa tctatcacag 120
 cctttagta ttgattgctc aatctataaa gagctcagtt tacagcatga ctgttagtaa 180

```
<210> 508
<211> 129
<212> DNA
<213> Homo sapien
```

```

      <400> 508
ctgttaaaag aacaaactta gcaatatata acagttnggt aacaggattt ttgactattc      60
actttggggag ttatatttttaa aaatccactt ttttactgag tcttactaca taccaggcac      120
tgtacttgg                                     129

```

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<220>  
<221> misc_feature  
<222> (1)...(422)  
<223> n = A,T,C or G
```

```
<210> 510
<211> 238
<212> DNA
<213> Homo sapien
```

<210>	511
<211>	254
<212>	DNA

<213> Homo sapien

<220>

<221> misc_feature

<222> (1)...(254)

<223> n = A,T,C or G

<400> 511

ccnattgatt	tgatggtaag	ggaggggatcg	ttgnggctcg	tctgttatgt	aaaggatgcg	60
tacggatggg	agggcgatga	ggactaggat	gatggcgggc	aggatagttc	agacggtttc	120
tatttcctga	gcgtctgaga	tgtagtatt	agttagtttt	gttgtaagng	ttaggaaaag	180
ggcatacagg	actaggaagc	acgataagga	aaatgactat	gagggcgnga	tcatgaaagg	240
tgataagctc	ttct					254

<210> 512

<211> 269

<212> DNA

<213> Homo sapien

<220>

<221> misc_feature

<222> (1)...(269)

<223> n = A,T,C or G

<400> 512

cctacctgta	aactacagta	ctttatatat	ctatgggntt	aataaaaana	aaatccacaa	60
atcttaaaaa	ggaactttaa	atgcagggct	atattgaatt	ggnaaaactgc	aacacaaact	120
ggcgcaacat	aggtaaata	ataccaatct	cactctatgt	gatgcaagca	tgctaactttc	180
ccactaattt	aaattacttt	caaccactat	gagccagaat	gcatgcctga	accttaaaact	240
gcactttaaa	aagtaacatc	ttggcctaa				269

<210> 513

<211> 266

<212> DNA

<213> Homo sapien

<220>

<221> misc_feature

<222> (1)...(266)

<223> n = A,T,C or G

<400> 513

ggaggggggt	tgttaggggg	tcggaggaga	aggntgggga	acagctaaat	aggttggtgt	60
tgatttggtt	aaaaaatant	agggggatga	tgctaataat	taggctgtgg	gtggttggtg	120
tgattcaaat	tatgtgnttt	ttggagagnc	atgncantgg	tagtaatata	attgttgaga	180
cgattagttt	tagcattgga	gtaggttttag	gttatgnacc	gtactctagg	ccatatgtgt	240
tgganattga	nactagtagg	gctagg				266

<210> 514

<211> 271

<212> DNA

<213> Homo sapien

<220>

<221> misc_feature
 <222> (1)...(271)
 <223> n = A,T,C or G

<400> 514
 acatgcaana aatcgagaat cttaaaaaac annacgaanc tgccttgga nncttactgg 60
 nntangatat ttatnttgcg gctgagatac ttgaacaact tcggatcnga antagacaan 120
 aangggnant tntatactgc nncagagggt acacagntca ttgtattaga gangaacana 180
 tgggtctggt gttcacacat tggggggaan atgggcgttn acangagagg nnganaaacn 240
 anganagcct ncttggttng cataanaaaa a 271

<210> 515
 <211> 328
 <212> DNA
 <213> Homo sapien

<220>
 <221> misc_feature
 <222> (1)...(328)
 <223> n = A,T,C or G

<400> 515
 ccaatgaggg gcaaagtgag cgncnagaag angttttgac tgaaataaat caaacacaaa 60
 aatntaagtt cacagtgaca gtttaaacaa aatccaaaca aactaacaac anaaacaccc 120
 cttgntttgc ctctagtggg aggtgggana acacaanctc gtccataaaa ttgactagta 180
 aaggggaaaa cccggtcatt tncctactct ttccangaaa tatctaatagc aagaaagaac 240
 ttctnctcat tatacngaag gaatttngaa aaatgatgta tttttggaac acctaantga 300
 aatactggaa cctgggcaag ttcaccac 328

<210> 516
 <211> 220
 <212> DNA
 <213> Homo sapien

<220>
 <221> misc_feature
 <222> (1)...(220)
 <223> n = A,T,C or G

<400> 516
 ncctnagttg aaggacccca tgtacatata gccagggga gcagtactag gntaactaga 60
 aggatctcat ccccatatgt gggctcattt caagtctatg gatgactacc ttcattgntg 120
 tgtgcgagat ggtttcaccc cttgaaaata tgggcacttc ancataanat agcnaaatct 180
 ttataatgat caatncatcc tacctccttt tacatgcatg 220

<210> 517
 <211> 296
 <212> DNA
 <213> Homo sapien

<400> 517
 tgcgatttct tccttggttg ttgctttggt ctgtgttcaa tccagagagc ttaaattgtc 60
 attatttttg gaagaaaacc tgtatttttg ttagtttaca atattatgaa atttcacttc 120
 aggagaaact gctgggcttc ctgtggcttt gttttcttag tttctttttc cgtgccgtgt 180

```
<210> 518
<211> 299
<212> DNA
<213> Homo sapien
```

```
<220>
<221> misc_feature
<222> (1) ... (299)
<223> n = A,T,C or G
```

```
<210> 519
<211> 464
<212> DNA
<213> Homo sapien
```

```
<220>
<221> misc_feature
<222> (1)...(464)
<223> n = A,T,C or G
```

```
<210> 520
<211> 221
<212> DNA
<213> Homo sapien
```

<210>	521
<211>	312
<212>	DNA

ccagacctgc	agaaaaactt	agcacagctc	aatctgctgt	tttgatggct	acaggggttta	60
tttggccaag	atactcactt	gtaactattc	caaaaaattg	gagtctgttt	gctgttaatt	120
tctttgtggg	ggcagcagga	gcctctcagc	tttttcgtat	ttggagatat	aaccaagaac	180
taaaagctaa	agcacacaaa	taaaagagtt	cctgatcacc	tgaacaatct	agatgtggac	240
aaaaccattg	ggacctagtt	tattatttgg	ttattgataa	agcaaagcta	actgtgtgtt	300

```

tagaaggcac tgtaactggt agctagttct tgattcaata agaaaaatgc agcaaacttt 360
taataacagt ctctctacat gacttaagga acttatctat ggatattagt aacatttttc 420
taccatttgt ccgtaataaa ccatacttgc tcaaaaaaaaa aaaaaacctt c 471

```

```

<210> 525
<211> 332
<212> DNA
<213> Homo sapien

```

```

<220>
<221> misc_feature
<222> (1)...(332)
<223> n = A,T,C or G

```

```

<400> 525
ccccnctgta ttccagcctg ggtgacccca tctcanggaa gaaaagttac cagatgtcgn 60
gggtaaaggt tggctctcaa gtggcctcat aagttgtctt gcattttaat tcaggggaatt 120
cattggacca ataggttaca ttttcgttcc ttttttgttt tggttcatct gttaagcagt 180
gggggcctaa ttactgctcc tttgtaaaaa cacattttcc caaagaacac tgaattaccg 240
ttcaaaactgg ttgttgatgg gtaataaggg ctgtttttgc tgccccaaaa gggcttaaca 300
atttaggcgg atagtttact taaaaaaaaa aa 332

```

```

<210> 526
<211> 440
<212> DNA
<213> Homo sapien

```

```

<220>
<221> misc_feature
<222> (1)...(440)
<223> n = A,T,C or G

```

```

<400> 526
ccaggttacc tcccctaaca gatgtggtgt tctganggggt tggttaagtg cccgaggaaa 60
ataggcctta actgttaaca tctacagaga agaaagcatg gtcacactgg caaggagtaa 120
gaagggattg ggtaaaagaa aatgggagag aaaagggaaa aaagtttttg caagacaatt 180
gttcctctgt aagaagctgc agggtgaaa ctttcctttc ttctattttt gtttttaatg 240
nctgtctctc tgatcagngg aaaagtgaag atttctagta tctagcacta acgtatgacc 300
caactttgag ggatcacaag ctagaacaag ttgaggattt aaaatcctgg ataattatat 360
acttaaagtt catgagcata aagctcactt gaccatgcag aaatgctggg aagcaggggtg 420
catggcatgg gaatacatct 440

```

```

<210> 527
<211> 124
<212> DNA
<213> Homo sapien

```

```

<220>
<221> misc_feature
<222> (1)...(124)
<223> n = A,T,C or G

```

```

<400> 527
tttccatag tctgttgggt gcataaatgn cttcttctga gaagtgtctg ttccctatcct 60

```

ttgccccctt tttgaggact taaatgttag acctaagacc ataaaaaccc tagaagaaaa 120
ccta 124

<210> 528
<211> 162
<212> DNA
<213> Homo sapien

<220>
<221> misc_feature
<222> (1)...(162)
<223> n = A,T,C or G

<400> 528
ctgcgggaga aatatgggga caagatgttg cgcangcaga aaggtgaccc acaagtctat 60
gaagaacttt tcagttactc ctgccccaaag ttctgtgcgc ctgtagtgcc caactatgat 120
aatgtgcacc ccaactacca caaagagccc ttctgcagc ag 162

<210> 529
<211> 409
<212> DNA
<213> Homo sapien

<220>
<221> misc_feature
<222> (1)...(409)
<223> n = A,T,C or G

<400> 529
cctttaaaat atagcttata aaatgtatac tatnngccag gagagctcac atttttctgc 60
agttttccag tggacctgcc tatggaatac tgtaaagaaa aatctgcaaa aatattccta 120
gcaattgaat cagtgccttt aaataaaaga agtggagagg ggcttggtta aattattctg 180
acaagttttc ttgctagtgg ttgccaaaat taaggatatt tgaagtgtcc tatcacccaa 240
atttggtttt aagaaaaagc tatattctgn gtctataggg tgaagccac actatctgtg 300
ctgcattctc aatgatacaa tacctatctg gaaactttcc tgttttgcca atgggtgcac 360
aaatctaaaa cattttatca caaagggtac ttgaatttaa atttctttt 409

<210> 530
<211> 325
<212> DNA
<213> Homo sapien

<220>
<221> misc_feature
<222> (1)...(325)
<223> n = A,T,C or G

<400> 530
ccgccagtgt gatggatata tgcagaattc gccctttcna gatttgngcc cgggcaggtc 60
catggctagg attatagata gttgggtggt tggggnaaat gagtgaggca ggagtccgag 120
gagggttagtt gtggcaataa aaatgattaa ggatactagt ataagagatc aggttcgtcc 180
tttagtggtt tgtatggcta tcatttggtt tgagggttagt ttgattagtc attgttgggt 240
ggtaattagt cggntgttga tganatattt ggagggtgggg atcaatagag ggggaaatag 300
aatgatcagt actgcggcgg gtagg 325

```
<220>  
<221> misc_feature  
<222> (1)...(173)  
<223> n = A,T,C or G
```

```
<210> 532
<211> 395
<212> DNA
<213> Homo sapien
```

<400>	532						
caggtcctac	tatgggtgtt	aaatTTTTTTT	ctctctctac	ngggTTTTTTT	cctagtgtcc		60
aaagagctgt	tctcttttg	actaacagtt	aaatttaca	ggggatttag	agggttctgt		120
gggcaaattt	aaagttgaac	taagattcta	tcttgga	ccagctatca	ccaggctcgg		180
taggtttgtc	gctctacct	ataaatcttc	ccactatttt	gtacataga	cgggtgtgct		240
cttttagctt	ttcttaggta	gctcgtctgg	tttcgggggt	cttagctttg	gctctccttg		300
caaagtattt	tctagttaat	tatttatgca	naaggtatag	gggntagtc	ttgctatatt		360
atgcttggn	ataatTTTTT	atctttccct	tgcgg				395

```
<220>  
<221> misc_feature  
<222> (1)...(290)  
<223> n = A,T,C or G
```

$$\begin{array}{ll} \langle 210 \rangle & 534 \\ \langle 211 \rangle & 334 \end{array}$$

<212> DNA
<213> Homo sapien

<220>
<221> misc_feature
<222> (1)...(334)
<223> n = A,T,C or G

<400> 534
ccgccagtgt gatggatatc tgcagaattc gcccttagcg agnnagccgg gcaggtccat 60
ggctaggttt atagatagtt ggggtggttg tggggnatga gtgaggcagg agtccgagga 120
ggttantttg tggcaataaa aatgattaag gatactagta taagagatca gggtcgtcct 180
ttagtggtgc gtatggctat catttgtttt gagggtagnt tgattagnca ttgttgggng 240
gtaattantc ggctgttgat ganatatattg gaggtgggga tcaatanagg gggaaatana 300
atgatcagtn ctgcggcngg tnnagacctn gccc 334

<210> 535
<211> 557
<212> DNA
<213> Homo sapien

<220>
<221> misc_feature
<222> (1)...(557)
<223> n = A,T,C or G

<400> 535
nccataagct tcagtgcgca aaaggtcaag gccagtgtta atttgttatt tcttaaataa 60
ctttcccttt cattttttaa ttataaattt aacttctaac atgttttatg gttaaaattg 120
tacttttttc ctttagcgac attcaaattc atcacaatca ctttgtgaaa ttgttcgcct 180
gagcagagac cagatgttac aaattcagaa cagtacagag cccgaccccc tgcttgccac 240
tctagaaaag tatgtgtaaa actctgttct tgttcttctt tcatattgat gctgttccat 300
gtgttaccat tgtgagtggg tggttaagtgt tccttatgtg ggaatcatgt gccttgaaaa 360
taaccttggg tgggtgagaa ggtagggaaa cctgcttctt ttatctcaag taaaagtttt 420
ggcagggtaa agaagataaa tgacatttat atctagactt ttgagttttc caattatttg 480
gtaaaaatgg gaaattctgt agaagccctt ccttaaaaat gggggaagtc catttnanaa 540
aattaactgg taggtca 557

<210> 536
<211> 372
<212> DNA
<213> Homo sapien

<220>
<221> misc_feature
<222> (1)...(372)
<223> n = A,T,C or G

<400> 536
gttccaacct tcattttctga aactgttcta gagcacngtg tctttctcgt agttcataac 60
ttacccttct agtctagaat tagaattaca ttatctgttt tactacttta ctagactgta 120
agtccttaga agataaggac tagggagttc atctctgtat tccaccagaa ggtacagtga 180
ctcatatcta gagtcttttag atgaaactta ctgagttgaa taacttaata tatttctgtt 240
ttcattccca agggaggcca tgtctggaga tagacctga atttaataaa ttttaggcac 300

```
<210> 537
<211> 284
<212> DNA
<213> Homo sapien
```

<400> 537

```
<210> 538
<211> 293
<212> DNA
<213> Homo sapien
```

<400> 538

```
<210> 539
<211> 468
<212> DNA
<213> Homo sapien
```

<220>

```
<221> misc_feature
<222> (1)...(468)
<223> n = A,T,C or G
```

<400> 539

$\langle 210 \rangle$	540
$\langle 211 \rangle$	397

<400> 543

```

cctactatgg gtgttaaatt ttttactctc tctacaaggt tttttcctag tgtccaaaga      60
gctgttcctc tttggactaa cagttaaatt tacaagggga ttttagagggt tctgtgggca     120
aatttaaagt tgaactaaga ttctatcttg ggcaaccagc tatcaccagg ctcggtagggt     180
ttgtcgccctc tacctataaa tcttcccact attttgctac atagacgggt gtgctctttt     240
agctgttctt aggtagctcg tctggtttcg ggggtcttag ctttggctct ccttgcaaag     300
ttattttctag ttaattcatt atgcagaagg tataggggtt agtccttgct atattatgct     360
tggttataat ttttcatctt tcccttgcgg tactatatct attgcgccag gtttcaattt     420
ctatcgcccta tactttatctt gggtaaattg tttggctaag                          460

```

```

<210> 544
<211> 116
<212> DNA
<213> Homo sapien

```

```

<220>
<221> misc_feature
<222> (1)...(116)
<223> n = A,T,C or G

```

```

<400> 544
ccgccagtgt gatggatctc tgcagaattc gccctttgga gngctnngcgc ccgggcagggt      60
ctgttttcagc agctcctcct tcttcttccc gcgangatct cgagccttga tcttgg       116

```

```

<210> 545
<211> 380
<212> DNA
<213> Homo sapien

```

```

<220>
<221> misc_feature
<222> (1)...(380)
<223> n = A,T,C or G

```

```

<400> 545
cgacggatcg atnagctnga tatcgaattc ggacgagcat ggcgtattgc tgcagatatg      60
gattcttcag aatgctccat gacaaatgta ctgacgggaa gncnatctaa aggaggcatt     120
gtnatgagag aaaggtctcg agctccagat aaagagagat acagagttct tgggaattgga     180
gttgcgaaaa cagtaagaca atcgattgtg gggaagcgtt ctttttagaga atctttggcc     240
ttcactccaa agcgtttgtt ttcattcaata ataagtagct cgtgccgaat tcctgcagcc     300
cgggggatcc actagttcta gagcggccgc caccgcggag gagctccagc ttttgttccc     360
tttagtgagg gtttaatttcg                          380

```

```

<210> 546
<211> 418
<212> DNA
<213> Homo sapien

```

```

<400> 546
ccaggggcaat taggcaggag aaggaaataa aggggtattca attaggaaaa gaggaagtca      60
aattgtccct gtttgccgat gacatgattg tataatctaga aaacccatt gtctcagccc     120
aaaaatctct taagctgata agcaacttca gcaaagtttc aggatacaaa atcaatgtac     180
aaaaatcaca agcattctta tacaccaata acagaccaac agagagccaa attatgagtg     240
aactcccatt cacaattgct tcagagaata aaatacctgg gaatccaact tacaagggat     300
gtgaaggacc tcttcaagga gaactacaaa cactgctca aggaaataaa agaggatata     360

```

```
<210> 547
<211> 172
<212> DNA
<213> Homo sapien
```

```
<210> 548
<211> 367
<212> DNA
<213> Homo sapien
```

<400>	548						
ggtctgactt	aagagaaaca	atggaaggca	agaggcagta	gaataatata	ttcaaaagat		60
gcaaaggaaa	aaaacctctc	agccacgaat	tccttatcca	gcaattattt	ttcaaaaatg		120
aaaataaacac	aaagacttag	ccagataaac	agaaacatta	actgaagttg	ttgctggcag		180
acctaccata	taaaaataaa	aaactctaaa	aaaattccta	tggctaaaag	caagttacag		240
aagacagtca	cttgaatcca	catttttaaaa	aaagcactga	tatacgtaat	attgacatta		300
taaaagacag	taaaaatgca	tttcttcttt	ataataaaatn	gcttattaaa	taacatgtgt		360
ataatgg							367

```
<210> 549
<211> 418
<212> DNA
<213> Homo sapien
```

<400> 549									
ccaa	atcaga	ac	tagagt	g	agcattctat	aaactcacct	ttgctttgat	ccttgaagat	60
caca	agtttt	g	tactgttg	g	aaatctctac	tctttcaaca	ctttaattaa	atggcattta	120
ga	atttcata	t	acttctggt	t	gttgtttcca	caatcttaaa	ctggatttag	aaatacttat	180
aat	gtaaaatg	c	aagagcttt	a	acttagtaa	ccgtatttcc	tattttttgt	tgtttttctt	240
ttg	ccagaat	t	tctgtttgt	t	ctacaataaa	gtccagcgaa	atacagtatt	tgggttaggtt	300
act	gttaac	a	ataaaat	t	atcatttgta	gagtttttac	ttaaccttcc	tattctctag	360
tct	ctataat	c	tttcaatga	a	gataaccag	ttacgaatat	ctcctatacc	atattagg	418

```
<210> 550
<211> 234
<212> DNA
<213> Homo sapien
```

```
<220>
<221> misc_feature
<222> (1)...(234)
<223> n = A,T,C or G
```

```

<400> 550
cctacccgcc gcagnactga tcattctatt tccccctcta ttgatcccca cctccaaata      60
tctcatcaac aaccgactaa ttaccaccca aactcacaa caaaactaac taataactaac      120
atctcagacg ctcaggaaat agaaaccgtc tgaactatcc tgcccggccat catcctagtc      180
ctcatcgccc tcccatccct acgcatcctt tacataacag acgagggtcaa cgat          234

```

```

<210> 551
<211> 542
<212> DNA
<213> Homo sapien

```

```

<220>
<221> misc_feature
<222> (1)...(542)
<223> n = A,T,C or G

```

```

<400> 551
caccctacc ccnntcctca taaaagttn ctcctctgga tcctcttttt cctcatgag      60
tgcccgggtg cccaagtcaa aaacctggga gtgatataaa ctccccacac atccagtcag      120
tcactcatca actctattga ttctgtctgc taaatatatn tcaattgtat taacttaaac      180
atatgcatan ggcactttct tcttcaactgc atttttgtgg gctgcactta cctttcaggt      240
aacgacaaca ctggccccctc ttgcccttct agtcagaagt gccaaaatga tgagagctag      300
ccatgacaaa cccacagcca acattacact gaatgtgcaa aactggaagg gcatccaaac      360
agaggagggg agagaggaat agacaggaag tcaaactgtc tctgtttaca gatgacatgt      420
ttctatatct ataaagcccc atagtcttgg ccccaaagct tcttctgctg ataaacttta      480
gcaaagtctt agcatacaaa atcaatgtgc aaaaattact aacagtccta tacatcaagt      540
ca                                         542

```

```

<210> 552
<211> 411
<212> DNA
<213> Homo sapien

```

```

<220>
<221> misc_feature
<222> (1)...(411)
<223> n = A,T,C or G

```

```

<400> 552
cctggntgac aaggaggtgc ctgtnatgtg aagatttgag gaaagagcat tccaggcagg      60
gggaaggcct gatgcaaagg gtctactgca ggcattagct gagcttattt aaagatcaga      120
atgaaggcca ttgtggctag aacagagtgg acaggaagga atgggtaccag gcaaagctga      180
agaagttggc aggattgagc tctcataant catggcaaag agttccattt tcattgtttg      240
acggaaataa attggaagggt cttaagtagg agaagatttg attagattta cattttacga      300
agaagcactc tggatgttat gtgaagaaat ggcctttgca gggcaagggt ggaaacaaag      360
agatcagtta ggaaattatt ggagtagctg aggattggat gaggggatgt g              411

```

```

<210> 553
<211> 631
<212> DNA
<213> Homo sapien

```

```

<220>

```

<221> misc_feature
 <222> (1)...(631)
 <223> n = A,T,C or G

<400> 553
 ccgggattag aactaaaaca agtgagatca cccctctaata ttttctgaa cttgggttaat 60
 aaaagtttat aagattttta tgaagcagcc actgtatgat attttaagca aatatgttat 120
 ttaaaatatt gatccttccc ttggaccacc ttcattgttag ttgggtatta taaataagag 180
 atacaaccat gaatatatta tgtttatata aaatcaatct gaacacaatt cataaagatt 240
 tctcttttat accttcctca ctggccccct ccacctgccc atagtcacca aattctgttt 300
 taaatcaatg acctaatgac aacaatgaag tttttataa atgtatttat gctgctagac 360
 tgtgggtcaa atgtttccat tttcaaatta tttanaattc ttatgagttt aaaatttgta 420
 aattttctaaa tccaatcatg taaaatgaaa ctggttgctcc attggagtag tctccaccc 480
 aaatatcaag atggctatat gctaaaaaga gaaaatatgg tcaagtctaa aatggctaata 540
 tgtcctatga tgctattatc atagactaac gacntttatc ttcaaaacac caaattgtct 600
 ttagaaaaat taatgtgatt acaggtagag g 631

<210> 554
 <211> 558
 <212> DNA
 <213> Homo sapien

<220>
 <221> misc_feature
 <222> (1)...(558)
 <223> n = A,T,C or G

<400> 554
 ccaggntagt ctccaactcc tgaccttagc tgatccaccc acctcggcct cccaaagtgc 60
 tgggattaca ggcattgagcc actgcgcccg gccaaacttg atatgcattt ttaaataagt 120
 taatacatta ttcattggttt agtctcatta tatattctat ggtccacttt gaaatttcat 180
 ctaacccaaaa tcatcttcat cctgcaattt gaggtttgga cacaatgggg attgatcagt 240
 aattttcttca tatgcccttt ctcaaggaaa tagtttccta tgaaaaaaa gtcctatggt 300
 ttcattgtaag ttctcttttt ggagaagaaa aggagacatt cttacttagc actctcagtt 360
 ttacaaaacg ctgccaacct taaaatttgt ctattgattc ccaaggcaca caaccaatag 420
 tctgtcaata acccggaata acatttcttt aaggccccag taactttcac atgtttgggt 480
 tccaatcttc acctagaatc ttgttaagaa aagtaaacca ttcactcctc tagaaactct 540
 aaggttgctt cttagggg 558

<210> 555
 <211> 212
 <212> DNA
 <213> Homo sapien

<400> 555
 ccagggtattt gcataatggc ttttcttctg ttgcctttgt tcctttgtgg cccagctaa 60
 ttgcctgaga gtgccactgt tagttttcaa ctctttctga tagaaaccct gtgtactaac 120
 atggaaatct taggtaatct gctttttcaa agcacaatgc agaatttatt ggcgggtggtg 180
 taactttaag aatatccgag aagccaccaa gg 212

<210> 556
 <211> 219
 <212> DNA
 <213> Homo sapien

<220>
 <221> misc_feature
 <222> (1)...(219)
 <223> n = A,T,C or G

<400> 556
 ccatgtgtct atctggagag aaggggaaac agcaagtgca aaggccctga gatggaacat 60
 atctggagaa ttcgaagaat ggtaagaagg ccagagtgga gcagaacaag tgtgggagag 120
 agttgtagga gatgagatca aaggctagga atgaagtgta aggccatgtc atgtgacctt 180
 gtatgtcctt gtaaggcttt tttttttttt tttnancct 219

<210> 557
 <211> 482
 <212> DNA
 <213> Homo sapien

<400> 557
 cctactatgg gtgttaaatt ttttactctc tctacaaggt tttttcctag tgtccaaaga 60
 gctgttcctc tttggactaa cagttaaatt tacaagggga tttagagggt tctgtgggca 120
 aatttaaagt tgaactaaga ttctatcttg gacaaccagc tatcaccagg ctcggtaggt 180
 ttgtcgctc tacctataaa tcttccact attttgctac atagacgggt gtgctctttt 240
 agctgttctt aggtagctcg tctggtttcg ggggtcttag ctttggtctt ccttgcaaag 300
 ttatttctag ttaattcatt atgcagaagg tataggggtt agtccttgct atattatgct 360
 tgggtataat ttttcatctt tccttgctg tactatatct attgcgccag gtttcaattt 420
 ccategccta tactttattt gggtaaatgg tttggctaag gttgtctggt agtaagggtg 480
 ag 482

<210> 558
 <211> 679
 <212> DNA
 <213> Homo sapien

<220>
 <221> misc_feature
 <222> (1)...(679)
 <223> n = A,T,C or G

<400> 558
 ctgtnaaaat tctgaaccta tccccaaaag aaaaaccgtg aaatacaagt tttaggaggt 60
 ggagcaaaga aaagccaagt tattttaaac caataaacac aagagacaat tctgctggag 120
 aatttacttt ctccaaaaca tcaaattggac tttaaagcag aagaccacat tttatgagaa 180
 agttatgtca ctgaaaagct tcatgtaaag tgactttgta aatggaatat ttttaaataa 240
 taaaaagaaa ataacttttc caggaatcct ttggagaggc tgataaccag atattaaatt 300
 atcaattttg ccaaagtgga ctttttaaaa atgtgttact tttaaaaact aacttgaaag 360
 aatttatgag gcaatctatc tgagtatggt tattgttgct ccattggctt tcaggatttt 420
 ggtcatttca ctgttaactc ttacatcaga gaataaagaa aagaaaatga aactttgtta 480
 ggaactggga tggaaaatgt agtcccagac agatctactg acctcgactg agtttcagaa 540
 atatcccagg attttggtta ttcatgcctt tcttttgtag ctttctttca aattagccaa 600
 ttaaagatac ccttcaatc accggtgaca tcagtacaac agtttttcaa cagttttctc 660
 tctcctgacc aaacagttt 679

<210> 559
 <211> 488

<212> DNA
<213> Homo sapien

<220>
<221> misc_feature
<222> (1)...(488)
<223> n = A,T,C or G

<400> 559
ccccactgta ctccagcctg ggtgacccca tctcaaagaa gaaaagttac cagatgtcat 60
gggtaaagggt tgggtcttcaa gtggcctcat aagttgtctt gcattttaat tcaggaatt 120
cattggacca ataggttaca ttttcgttcc ttttttgttt tggttcatct gttaagcagt 180
gggggcctaa ttactgctcc tttgtaaaaa cacattttcc caaagaacac tgaattaccg 240
ttcaaaactgg ttgttgatgg gtaacaagggt ctgtttttgc tgccccaaaa gggcctaaca 300
atttaggcgg atagtttact taaaaaaaaa aatcctttgg agacatactg aaaatgcaaa 360
ctagtttcta aattatcaat tccctacatg aanaagcagt ttgccanagt ttagtctcan 420
aaaatgactg gttggctcta tttaaatcan aacccaattt ctacgcacct gccgcgccgg 480
ccaagggc 488

<210> 560
<211> 602
<212> DNA
<213> Homo sapien

<220>
<221> misc_feature
<222> (1)...(602)
<223> n = A,T,C or G

<400> 560
cctanttaag aattccttgc cttagtgggt aacaaggact aaacacagac aatgggtgaa 60
acacagacgc taattcacat aacagagagt aggcaacctt aagaatgaat tgatgcagac 120
tcctatagaa ttctctgtgt atgactgggt tcttattttc tcctccttgt atgtagtga 180
aatttcatca ttatgaatag ttcttggat ctttttttaa agttgtgaat gcgagtgttt 240
ggctttgtaa tacaactttt tagtatccag aagataacca gtgctctacc aataaagatc 300
ttttgataca aagggtttta acttctgcca gttcttactc atttttttca ggttttttat 360
acatttctta aacaacacat acattatgta aaatataaga attaatgtac attctcaagg 420
ccagattcag tgacaaaatg cactaccgca atctagtaac acatttactc cttgctgcat 480
ataagtggcg tgtaagaaat acagggtata ttgttttgtg atccatgcag taaatgttca 540
caaatatcag gcaacaact agacgntctt cagctactaa aattaactgt cccagtcaca 600
aa 602

<210> 561
<211> 683
<212> DNA
<213> Homo sapien

<400> 561
gtctatTTTTT aaaaagaaag aaaaaaacca cttttttata gtccttagct ttgccatatg 60
cccgccttaa gtggaaggaa agttaatcac ttaactatgt ttataaaaaa gaaaaaagggt 120
cttggaatgc tattactgtt cacacaaagt atgattctgt ttgaataagg caaatgctcc 180
tttttttaaa aaaagacatt actgtaatat caaaaaccgt ggcagtttgt atacaactct 240
gggcttgatt ttttttaaaa aaacagaatg aattgatgtc ttattttata aatgttctat 300
atttattagg agaaaacttt atattgcctt ttttatcaat catgtaacag gcttatagct 360

```

ttccaacaga gctgcttgcc aaacaat ttttgtttat taaacagtgc tgaaacaaac 420
aggatcagca tttacttaag atgttaagaa tgaggacttt taatcagccg aaccaagata 480
ttgttacctg tatgcattcc caaagtctag atgctcagta tgttcagtca tatctttcag 540
aatcagtga cccgattacc tttttttggt attcactcta catctgcca cctagttcac 600
cttggttttg tgtctgctgt agaaggggaa cataacttgg ttaaaccgta gggattatca 660
ttgtatacat gctgtgaaca tgt 683

```

```

<210> 562
<211> 420
<212> DNA
<213> Homo sapien

```

```

<400> 562
gcactttttt tccagtaagg attcatctct tgcctccta tatggtcatt atattttata 60
ttttacatat ttataaacat gacatatgta tttatgttcc acaaagggct ttgaatagaa 120
tttacacata gagttccctg gggtgatgtg tttatcaaaa tggaagataa agtgaattaa 180
ttactttaat atttaacact attgaataga aataatttcc ccaatattgc ttcattgattt 240
agacagtcta ttaaattgtt aagcaaggca ctagactaag tttattaaga caaattttgg 300
aatatgtgca gaaatatgac ctggctaata gtacagagtc aaagctgggt gaatgggtgtt 360
atatagtgga ttcagattga tgtggcagtg gtggttacac taggggcact aaggttatcc 420

```

```

<210> 563
<211> 482
<212> DNA
<213> Homo sapien

```

```

<400> 563
ctccacctta ctaccagaca accttagcca aaccattttac ccaaataaag tataggcgat 60
agaaattgaa acctggcgca atagatatag taccgcaagg gaaagatgaa aaattataac 120
caagcataat atagcaagga ctaaccctta taccttctgc ataataaatt aactagaaat 180
aactttgcaa ggagagccaa agctaagacc cccgaaacca gacgagctac ctaagaacag 240
ctaaaagagc acaccctgt atgtagcaaa atagtgggaa gattttatagg tagaggcgac 300
aaacctaccg ggcttggtga tagctggttg tccaagatag aatcttagtt caactttaac 360
tttggccaca gaaccctcta aatccccttg taaatttaac tgttagtcca aagaggaaca 420
gctctttgga cactaggaaa aaaccttgta gagagagtaa aaaatttaac acccatagta 480
gg 482

```

```

<210> 564
<211> 302
<212> DNA
<213> Homo sapien

```

```

<400> 564
ctggaagtga aggtactaat atacaaatgg ctcttgtttc tgaatatgtg atataatttg 60
tgaatctttg gaaactgaat tttttctatg gagtgcaaat atagaagggt tattttacaa 120
tgtttggtgt gaaaagaatt cactttgtaa acaactatta aggctggaag tttagtgaag 180
gtgcatagtt ttgaaagcta cacagggtgaa aaatcaaact tattgtttgt aattttgctg 240
ttacatgtta agttactttg acagcaattt tctaatagata atgtgattta tgatttaaaa 300
gg 302

```

```

<210> 565
<211> 554
<212> DNA
<213> Homo sapien

```

<220>
 <221> misc_feature
 <222> (1)...(554)
 <223> n = A,T,C or G

```

<400> 565
ccanngtgac atcatggcaa tacagcaaga attctgnnat ttatttagaa gcctcaagga      60
gaaggatcct ggagcccctg aatgagagtt tcttctccat gcctctcccc agtcaaaaata    120
catggaaata ttcatagaag cattgtaccc agcatgataa ggaaggatgg agaattggttc    180
cttatatctc tgttcacaag acatcaacac tcttaagtaa ctgtatgaaa taaattctctc    240
gctgaaagca aataaaccat ctgaaaggtc ttctgggttac ttacacagat ttcctagaga    300
atctgaaatc agcctaacag ggaagattaa tttttaaatg aatccaagtt aatgaaagca    360
aagaactctt atacagaaat acattttcct attataaagc aggactacct tccctaattt    420
ctgatatacc taggacaatt tgaatgggca ttgaaattct tttgggttgaa ttacgcaaac    480
aagcaaagga aaagtctcaa ttattattgg aaaatttggg gagagattat tatctcttga    540
tctcctagtn natt                                                    554

```

<210> 566
 <211> 631
 <212> DNA
 <213> Homo sapien

<220>
 <221> misc_feature
 <222> (1)...(631)
 <223> n = A,T,C or G

```

<400> 566
ncgaagctgt gaanncattc acacggaatc tgganggtat tactgtaact tcttataata      60
cataatataa aagtttttga aagatataga cacaattaac ccctaaacaa cacactatct    120
gattctcaaa agcaatggct atttaacaag atgtaaaagg acaataacat atcaaagaac    180
tttcacacac ctaaagatag catttagcag caagttagtc agacaaaaca aacataaata    240
tcttcacatt tcctatgttt gtttttaact ttacttcata aagccactga taattgaggt    300
ttcttttcaag tataagattt ctaaaattaa aaactgtttt tgacatatatt ttataaagaa    360
ataaaaagca aaacgcaatc caactattta tatgagtcct tcttctccaa cagctttaga    420
tgtttttctg agtacttttt acacagaata tttttattaa aatcagttct aattcattta    480
tgcagattag gggaaaatga ttcataataa attaacctta aaattacctt ctatctgctt    540
ctacctctat ccccccata ccaccaaatc tgttgctaca gtgaactgta gccaatgtct    600
gtttgagggg gcccaaagca tctggtaatc t                                                    631

```

<210> 567
 <211> 510
 <212> DNA
 <213> Homo sapien

<220>
 <221> misc_feature
 <222> (1)...(510)
 <223> n = A,T,C or G

```

<400> 567
cctatnatag cttctctagc tatcatactc caatcagcna aaaatgagaa aatgttgaga      60
aatagaagat aattcctcat ttaaggncac cttctanaat ttgtgcttaa nantctgttt    120

```

```

tcttctcatg ggccagcact tcggcaactg ggaaaaatta nngtacagg gatctaggna 180
atactgttta tttagagcaat aatatattgn gctaacgttc aggcataccta ttactgagaa 240
ataagggaaa atgagtgtaa agtacaacta agagtctcgg ctacaggga aaataccatc 300
agttaaatat ccatagtcct agagcattta tgtaaaactg caatttgaat cctgcaatac 360
attttggtt tttcctcagt gataccatgt gtgggaagtt gttctgtcaa ggtgggtcgg 420
ataatttgcc ctggaaagga cggatagtga ctttctgac atgtaaaaca tttgatcctg 480
aagacacaag tcaagaaata ggcattggtg 510

```

```

<210> 568
<211> 180
<212> DNA
<213> Homo sapien

```

```

<220>
<221> misc_feature
<222> (1)...(180)
<223> n = A,T,C or G

```

```

<400> 568
ttaatntgac ncacgcttat gcggaggaga atgntttcat gttacttata ctaacattag 60
ttcttctata gggatgata ttgggtccaat tgggtgtgag gagttcagtt atatgtttgg 120
gatttttttag gtatgggtg ttgagcttga acgctttctt aattgggtggc tgcttttagg 180

```

```

<210> 569
<211> 237
<212> DNA
<213> Homo sapien

```

```

<400> 569
ccaattgatt tgatggtaag ggagggatcg ttgacctcgt ctgttatgta aaggatgcgt 60
agggatggga gggcgatgag gactaggatg atggcgggca ggatagttca gacggtttct 120
atttcctgag cgtctgagat gttagtatta gttagttttg ttgtgagtgt caggaaaagg 180
gcatacagga ctaggaagca gataaggaaa atgactatga gggcgtgatc atgaaag 237

```

```

<210> 570
<211> 352
<212> DNA
<213> Homo sapien

```

```

<400> 570
ctgtctctcc atttagagcc ccagttggtc ctgacctctt acaaatttgg tgttttccact 60
ttgatgttta tgaaccgatt gcattaaaaa tgcaggataa tgattcaggg ttagagaaac 120
tattatttat acaaattgtg ttaacacctc atcattttta attggctgtg ctaataatgc 180
tcattgtgct cttcaggggt atgtgtgtgt gtgtgtgtgt gttttgcctg aatctgcaac 240
ctacatttgc tctggcagta tgttgagtat atgctagaat agaattggacc taggcaactc 300
taaggctcta caactaaata cacttactta ggaaacctcc taaataagta gg 352

```

```

<210> 571
<211> 402
<212> DNA
<213> Homo sapien

```

```

<400> 571
ctgattttta caataactac tgtgttctct gcaatagtgt gttctgatta gaaatgacca 60

```

```

atattatact aagaaaagat acgactttat tttctggtag atagaaataa atagctatat 120
ccatgtactg tagtttttct tcaacatcaa tgttcattgt aatgttactg atcatgcatt 180
gttgagggtg tctgaatgtt ctgacattaa cagttttcca tgaaaacgtt ttattgtgtt 240
tttaatttat ttattaagat ggattctcag atatttatat ttttatttta tttgtttcta 300
ccttgaggtc ttttgacatg tggaaagtga atttgaatga aaaatttaag cattgtttgc 360
ttattgttcc aagacattgt caataaaagc atttaagttg aa 402

```

```

<210> 572
<211> 70
<212> DNA
<213> Homo sapien

```

```

<220>
<221> misc_feature
<222> (1)...(70)
<223> n = A,T,C or G

```

```

<400> 572
tggatccgag ctcggtacca agcttggcgt aatcatggtc atagctgttt cctgtgntcg 60
ttttacaacg 70

```

```

<210> 573
<211> 423
<212> DNA
<213> Homo sapien

```

```

<400> 573
ccaatggttt cttagtgaag gagtacctta gctctgaatg caatgccctc agaaagatat 60
cattcataga gacatacaaa gcacatggca acatgacatt ggaatacacg attctgagca 120
tcttcattca tgaccaacct ggctatagat ttcagatgtc ctcttggctc gaaggatata 180
tgggatatcc atgctcactt gcattccttt ccctttaatt tcattttcta agtccttctt 240
gtattgtttc taaaagaaca gaaaataatc ttggagcttt gcttaagctt taatagcgat 300
gttgaaattt acatgtttga atctcaaagc caccatgtg gaaagaaaac ttatgctctt 360
tccagctatg attcacggca tttattttta actttgtatc ttgctgctgt cttacctggc 420
tgg 423

```

```

<210> 574
<211> 129
<212> DNA
<213> Homo sapien

```

```

<400> 574
ctgttaaaag aacaaactta gcaatatata acagtttgct aacaggattt ttgactattc 60
actttgcgag ttatttttta aaatccactt ttttactgag tcttactaca taccaggcac 120
tgtacttgg 129

```

```

<210> 575
<211> 684
<212> DNA
<213> Homo sapien

```

```

<220>
<221> misc_feature
<222> (1)...(684)

```

<223> n = A,T,C or G

<400> 575

ccagatntga	cttttcaaaa	ctactcacat	tgtgaaaaan	gcaggaacaa	atctagtttc	60
aagttcagca	tgccgttccc	tgtttaattc	ataaaacaca	actggcagaa	gtattacttg	120
aagcaaaaca	aaagtaacgt	gggaacttgc	ttatttgcta	agccacaatg	tatttttcca	180
ggaatagcat	aaatttgcca	tctttcttgc	gtctatggaa	aaggggttta	gaattgtttc	240
actaaaaatt	aaatttctat	attgtcaaac	atgattgtat	actcaaattt	taaaatgtga	300
agggaacact	tactaagcat	ttcctgggta	tgccactata	ttaagtccta	gtaatatgat	360
atagtttatt	tcaatttttt	ttcaactcat	acttccttta	aaatagcact	gaccaaaga	420
aagttaacat	gagcttcatg	tacaattttt	aatctttttg	cagaaaaata	aactgagaaa	480
ggctaaaatt	gttttattta	agccactata	ccaagacata	ttgatttcac	caatataaaa	540
attgagatag	tttacatttt	ttggtacatc	tttaaaatct	ggtatgtatt	tttatactga	600
cagcacatct	caatttggtg	aagctacatt	tccagggctc	aatagtcacc	atgaatctca	660
attgtaatca	aagagggttg	cctg				684

<210> 576

<211> 134

<212> DNA

<213> Homo sapien

<400> 576

ccttatttct	cttgtccttt	cgtacagggg	ggaatttgaa	gtagatagaa	accgacctgg	60
attactccgg	tctgaactca	gatcacgtag	gactttaatc	gttgaacaaa	cgaaccttta	120
atagcggctg	cacc					134

<210> 577

<211> 133

<212> DNA

<213> Homo sapien

<220>

<221> misc_feature

<222> (1)...(133)

<223> n = A,T,C or G

<400> 577

ctgtctctcc	attnagaagc	cccantnggt	cctnacctct	tacaaatttg	gtgttttcac	60
tttgatgttt	atgaaccgat	tgcatataaa	atgcaggata	atgattcagg	gttaganaaa	120
ctattattta	tac					133

<210> 578

<211> 200

<212> DNA

<213> Homo sapien

<400> 578

cctcaaatct	atcttcaaa	gtgaccagc	aatcagtgtc	aatgccttta	ctgtagttaa	60
cctggtaatt	tcattcttta	gtctctccaa	gaaaatctga	agtgtattag	gcaagtcaga	120
acccaaattg	tctccaaggt	tgcaataaat	ttgtcccata	caggaaatag	ccctttcctt	180
gacttcctga	tcaatgtcag					200

<210> 579

<211> 402

<212> DNA

<213> Homo sapien

<400> 579

ctgatttttaa	caataactac	tgtgttctctg	gcaatagtgt	gttctgatta	gaaatgacca	60
atattatact	aagaaaagat	acgactttat	tttctggtag	atagaaataa	atagctatat	120
ccatgtactg	tagtttttct	tcaacatcaa	tggttcattgt	aatgttactg	atcatgcatt	180
gttgagggtg	tctgaatggt	ctgacattaa	cagttttcca	tgaaaacggt	ttattgtgtt	240
tttaatttat	ttattaagat	ggattctcag	atattttatat	ttttatttta	tttgtttcta	300
ccttgagggtc	ttttgacatg	tggaaagtga	atttgaatga	aaaatttaag	cattgtttgc	360
ttattgttcc	aagacattgt	caataaaagc	atttaagttg	aa		402

<210> 580

<211> 245

<212> DNA

<213> Homo sapien

<220>

<221> misc_feature

<222> (1)...(245)

<223> n = A,T,C or G

<400> 580

ccaattgatt	tgatggtaag	ggagggatcg	ttgacctcgt	ctgttatgta	aaggatgcgt	60
agggatggga	gggcgatgan	gactaagatg	atggcgggca	ggatagttca	gacngtttct	120
atttcctgag	cgtctgagat	gttagtatta	gttagttttg	ttgtgagtgt	taggaaaagg	180
gcatacagga	ctaggaagca	gataaagaaa	atgactntta	gggcgtgatc	atnaaanggg	240
ataaa						245

<210> 581

<211> 294

<212> DNA

<213> Homo sapien

<400> 581

tgcagcgcaa	gtaggtctac	aagacgctac	ttccccctatc	atagaagagc	ttatcacctt	60
tcattgatcac	gccctcatag	tcatttttct	tatctgcttc	ctagtcctgt	atgccctttt	120
cctaacactc	acaacaaaac	taactaatac	taacatctca	gacgctcagg	aaatagaaac	180
cgtctgaact	atcctgcccc	ccatcatcct	agtcctcatc	gccctcccat	ccctacgcct	240
cctttacata	acagacgagg	tcaacgatcc	ctccccttacc	atcaaatcaa	ttgg	294

<210> 582

<211> 230

<212> DNA

<213> Homo sapien

<400> 582

gaggtegcgc	tcattagtcac	tttccttctc	tgcttcctag	tcctgtatgc	ccttttctcta	60
acactcacia	caaaactaac	taataactaac	atctcagacg	ctcaggaaat	agaaacgcgc	120
tgaactatcc	tgcccgccat	cctcctagtc	ctcatcgccc	tcccatccct	acgcctcctt	180
tacataacag	acgagggtcaa	cgatccctcc	cttaccatca	aatcaattgg		230

<210> 583

<211> 481

<213> Homo sapien

[illegible]

<213> Homo sapien

ccaattaaga	gctaaattta	caaaataatc	tctatcagga	ggctttaagg	tttaatgtct	60
ctaaagtcct	tatggatata	agaggcttga	atgtactgaa	ttcaaatttg	gtttttaaat	120
gttataatag	tttaggcccc	agagccacat	atttctgtct	aagaatagaa	agcatagcta	180
gctgcccaca	cagaatattc	atatagaggt	gggggggcaag	aacaaaattt	attcatttga	240
tacatagaaa	tgggactact	tagaatagac	tcataataga	aagcatcatc	tggttttctca	300
tctcag						306

<213> Homo sapien

ccagaatggt	acagagtgga	gggtgttctg	ctaattgactt	cagagaagta	tttaagaaaa	60
acatagaaaa	acgtgtgcgg	agtttgccag	aaatagatgg	cttgagcaaa	gagacgggtgt	120
tgagctcatg	gatagccaaa	tatgatgcca	tttacagagg	tgaagaggac	ttgtgcaaac	180
agccaaatag	aatggcccta	agtgcagtgt	ctgaacttat	tctgagcaag	gaacaactct	240
atgaaatgtt	tcagcagatt	ctgggtatta	aaaaactaga	acaccagctc	ctttataatg	300
catgtcag						308

<213> Homo sapien

<223> n = A, T, C or G

cctgtctttg	aatggatgaa	atagggttaat	aaaaaacatc	actgttttaa	aactagaaca	60
ctgaaaaatt	ctaggaaagc	ttattttccc	ttatatTTTT	atggnaCTTT	caacacttna	120
caacactatt	tnaattaann	tttnttctag	agtttatann	atatcagtac	attctttttct	180

gtggatgcaa taatatagaa tcttattnca aatcttactg gcaggntctn ttaaattctt 240
 caacggntgn catagtgatt aacccaaaatt agttatgatt tctgcctatc tgtgtgagaa 300
 cttacagggg aaattgttct aaacctgagg aacatgaagt aactgtactg cacactccaa 360
 atgatgacag tcattttata tcaccttcaa ttaccaaca gcttttaata gtctgg 416

<210> 587
 <211> 382
 <212> DNA
 <213> Homo sapien

<400> 587
 cctactatgg gtgttaaatt ttttactctc tctacaaggt tttttcctag tgtccaaaga 60
 gctgttcctc tttggactaa cagttaaatt tacaagggga tttagagggt tctgtgggca 120
 aattttaaagt tgaactaaga ttctatcttg gacaaccagc tatcaccagg ctcggtagggt 180
 ttgtcgctc tacctataaa tcttcccact attttgetac atagacgggt gtgctctttt 240
 agctgttctt aggtagctcg tctggtttcg ggggtcttag ctttggctct ccttgcaaag 300
 ttatttctag ttaattcatt atgcagaagg tataggggtt agtccttgct atattatgct 360
 tggttataat ttttcatctt tc 382

<210> 588
 <211> 307
 <212> DNA
 <213> Homo sapien

<400> 588
 cctactcttc tccgtccatt gtactatctg cccgtgggtg ggatggcagt aggatcatat 60
 ttgatgactt ccgagaagca tattattggc ttctgcataa tactccagag gatgcgaagg 120
 tcatgtcctg gtgggattat ggctatcaga ttacagctat ggcaaaccga acaatttttag 180
 tggacaataa cacatggact aataccata tttctcgagt agggcaggca atggcgcca 240
 cagaggaaaa agcctatgag atcatgaggg agctcgatgt cagctatgtg ctggtcattt 300
 ttggagg 307

<210> 589
 <211> 89
 <212> DNA
 <213> Homo sapien

<400> 589
 cctgggtgat tgaggatgca atgagctgtg attgtgccac cacactccag cctgggcaat 60
 acagcaagac tgtctcaaaa aaaaaaaaa 89

<210> 590
 <211> 456
 <212> DNA
 <213> Homo sapien

<400> 590
 cctcagttct tgatttgtgt tgacggggcg tcaccatgaa ggagcccatt tagtataaag 60
 cttccaacct tttctcttaa tegtttcttt aatcttttaa accatcttca agtgcatagg 120
 ggagtttccg atgccagagg atgaaagcaa gtgctctctc caccctctcc tcccagagtg 180
 aaaacaaatc cttttgtctga tacttgtttc aaaagcatcc attgtaaagc ttctcagtga 240
 cacaaaatac tgagaggtaa ctttttatca atcaaaccac ataccccaat ttaacacctt 300
 tcaatgctct gaattcaact gacagactaa aggggtgttc ctgtaacagt ctgaaatatt 360
 aagtgttttt tttgttttgt ttttaaactc tatttcagaa aacttctct tggggtagga 420

aagtacacat gaagcagcaa agtaacgaag aaaaac

456

<210> 591
<211> 289
<212> DNA
<213> Homo sapien

<400> 591
ccaattgatt tgatggtaag ggagggatcg ttgacctcgt ctgttatgta aaggatgcgt 60
agggatggga gggcgatgag gactaggatg atggcgggca ggatagttca gacggtttct 120
atttcctgag cgtctgagat gttagtatta gttagttttg ttgtgagtgt taggaaaagg 180
gcatacagga ctaggaaagca gataaggaaa atgactatga gggcgtgatc atgaaagggtg 240
ataagctctt ctatgatagg ggaagtagcg tcttgtagac ctacttgcg 289

<210> 592
<211> 435
<212> DNA
<213> Homo sapien

<220>
<221> misc_feature
<222> (1)...(435)
<223> n = A,T,C or G

<400> 592
cgcgttagat gcgccttttc cggcctgtgc gtctgctctg gttcctctca ggcagcaaag 60
ctggggaagg aagctcaggc aggagcctcc ccgacaccac agcggcacia gcagcagcta 120
aagcaccgca ctttgctctg ctaacctttt acttaaatga ggttttgcca aatccacatc 180
tggaaccgca tcacacccat ttgcaaggat gtttggtctt tgatgaaact gcatctctac 240
tgcacatgan ggctttcatt gtaggacaag aggagagtgc gtttatTTTT gtaactgttt 300
tacetgttcc gattanttaa tcggnagctt atgtcatatt ctatgcctgt tgtcttctaa 360
tctctcctta ctaaaacatt acttcaaatt tnaattgacc cttgtttata atttatttaa 420
cgggatttgn gtgtc 435

<210> 593
<211> 633
<212> DNA
<213> Homo sapien

<220>
<221> misc_feature
<222> (1)...(633)
<223> n = A,T,C or G

<400> 593
ctgttttagtc agataattgt gtccgaattg attangaaaa taatagacca gccataaagc 60
agcataaaat attatgaaac tattccagaa gttcagtaat atctttggga cctgtctata 120
goccaaagttt tgtgaatact tttgtagtta aaaaaaattt ttactttacc agggcattgc 180
aattcttttc catcagtga tttcattcta cagacttttc agagcatctc ataatacagtc 240
aacaatatcta tttcaaatgt gtttggttact aagcaacggt tgctaagagc ttctgtaatt 300
aagatgaaag ttccaaggta acaatgccca aacacagcac cattttcacc attttctgat 360
aatgcaggag taggatggct aaaagtgaag gaagaatcta ctctatggaa agcatggcac 420
ctgaaatttc tgaagatatt ggctgtcctc tagcttatat gagagagagt gtttgtgctt 480
tactaatcaa ccagtcattt ttttcttgtg tggctgaaat gtacattcca gacatgaaca 540

ggtagagtat gtgttggggg caggtttata ctgcatgggt gtgctgagac agggccacgt 600
 ggtgatgtaa atgatgctgn ctgacacgtg cag 633

<210> 594
 <211> 501
 <212> DNA
 <213> Homo sapien

 <220>
 <221> misc_feature
 <222> (1)...(501)
 <223> n = A,T,C or G

<400> 594
 cctttacaag atgctggtac cttgatcttg gacngggcag gctccaagat ggaaagaaag 60
 tgagcatctg cttttttaggg attatccagt ctatactact ctgttctagc cacacaaaac 120
 aggttaagac agaaattggt accaagagtg ggggtgttact acagcaaata cctgaaaatg 180
 tagaagaggc tttgaaatgt ggtaattgga agaagctggg agaatttgga ggagtaggct 240
 agaaaatgtc tgtattttca tgaatggagc attaagaata attccgggtga ggccataggg 300
 aaagtctaaa actttttcaga aattatgtaa gcgattgtga ttagtagggtt ggtagaaata 360
 tagacagtaa aagcaattct gatgtgggtt cagaggaaaa tgaaaaatat tagaaactga 420
 aggaaggggc atccttgcta taaactggca aagaacttgg ctgaaatgtc tccatgtcca 480
 agagatttat ggcagaaatg t 501

<210> 595
 <211> 383
 <212> DNA
 <213> Homo sapien

<400> 595
 ctggtcacca tcatcccttt aatcaactca cacctgttta aagagtgttt ctgatttgac 60
 cttcatccct tagtttactg gcgttaaaaa aagtctcagc aattttcatt atttctcgtg 120
 ggtctcatta tcaaaccctt acttatttcg gcataatttc tctgggcttc ttctagtttc 180
 tgccttacia gcaatgctgt tctgtaaatt tattgaaacc tctggaacat ttcaccttta 240
 gagatggagg atggaaggat tgggtaccaga agagggctaa gatacgtttt ctgtcttgag 300
 ctgaaagcac agtctactct ccttcgtttt gtcgatgaga aagttgaggc cagagggggag 360
 gtgacatgtt tagagtcacc cag 383

<210> 596
 <211> 266
 <212> DNA
 <213> Homo sapien

<400> 596
 ccattggctag gtttatagat agttgggtgg ttggggtaaa tgagtgaggc aggagtccga 60
 ggagggttagt tgtggcaata aaaatgatta aggatactag tataagagat cagggttcgtc 120
 ctttagtggt gtgtatggct atcatttggt ttgaggttag tttgattagt cattgttggg 180
 tggtaattag tcggttggtg atgagatatt tggaggtggg gatcaataga gggggaaata 240
 gaatgatcag tactgcggcg ggtagg 266

<210> 597
 <211> 383
 <212> DNA
 <213> Homo sapien

<220>
 <221> misc_feature
 <222> (1)...(383)
 <223> n = A,T,C or G

<400> 597
 ctggtcacca tcatcccttt aatcaactca caccngttta aagagtgttt ctgatttgac 60
 cttcatccct tagtttactg gcgttaaaaa aagtctcagc aattttcatt atttctcgtg 120
 ggtctcatta tcaaaccttt acttatttgc gcataatttc tctgggcttc ttctagtttc 180
 tgccttacia gcaatgctgt tctgtaaatt tattgaaacc tctggaacat ttcaccttta 240
 gagatggagg atggaaggat tgggtaccaga agaggggctaa gatacgtttt ctgtcttgag 300
 ctgaaagcac agtctactct ccttcgtttt gtcgatgaga aagttgaggc cagaggggag 360
 gtgacatgtt tagagtcacc cag 383

<210> 598
 <211> 266
 <212> DNA
 <213> Homo sapien

<400> 598
 ccatggctag gtttatagat agttgggtgg ttggtgtaaa tgagtgaggc aggagtccga 60
 ggagggttagt tgtggcaata aaaatgatta aggatactag tataagagat caggttcgtc 120
 ctttagtggt gtgtatggct atcatttgtt ttgagggttag ttgattagat cattgttggg 180
 tggttaattag tcggttggtg atgagatatt tggaggtggg gatcaataga gggggaaata 240
 gaatgatcag tactgcggcg ggtagg 266

<210> 599
 <211> 294
 <212> DNA
 <213> Homo sapien

<220>
 <221> misc_feature
 <222> (1)...(294)
 <223> n = A,T,C or G

<400> 599
 ccaattgatt tgatggtaag ggagggatcg ttgaccacgt ctgttatgta aaggatgcgt 60
 agggatggga gggcgatgag gactaggatg atggcgggca ggatagttca gacggtttct 120
 atttcctgag cgtctgagat gttagtatta gttagttttg ttgtgagtgt taggaaaagg 180
 gcatacagga ctaggaagca nataaggaaa atgactatga gggcgtgatc atgaaagggtg 240
 ataagctctt ctatgatagg ggaagtagcg tcttgtagac ctacttgccg tgca 294

<210> 600
 <211> 213
 <212> DNA
 <213> Homo sapien

<400> 600
 agatattggg ctgttaattg tcagttcagt gttttaatct gacgcaggct tatgcggagg 60
 agaatgtttt catgttactt atactaacat tagttcttct atagggtgat agattgggtcc 120
 aattgggtgt gaggagtcca gttatatgtt tgggattttt taggtagtgg gtgttgagct 180
 tgaacgcttt cttaattggg ggctgccttt agg 213

<210> 601
 <211> 471
 <212> DNA
 <213> Homo sapien

<220>
 <221> misc_feature
 <222> (1)...(471)
 <223> n = A,T,C or G

<400> 601
 ncctactatg ggtgttaaatt tttttactct ctctacaagg ttttttccta gtgtccaaag 60
 agctgttcct ctttggacta acagttaaat ttacaagggg atttagaggg ttctgtgggc 120
 aaatttaaag ttgaactaag attctatctt ggacaaccag ctatcaccag gctcggtagg 180
 tttgtgcct ctacctataa atcttccac tattttgcta catagacggg tgtgctcttt 240
 tagctgttct taggtagctc gtctggttcc ggggggtctta gctttggctc tccttgcaaa 300
 gttatttcta gttaattcat tatgcagaag gtataggggt tagtccttgc tatattatgc 360
 ttgggtataa tttttcatct ttcccttgcg gtactatatc tattgcgccg gggttcaatt 420
 tctatgcct atactttatt tgggtaaatt gtttggtctaa ggttgctctgg t 471

<210> 602
 <211> 482
 <212> DNA
 <213> Homo sapien

<220>
 <221> misc_feature
 <222> (1)...(482)
 <223> n = A,T,C or G

<400> 602
 tgagcataca gcaataaaaa taacataatt tntatgtgta caatatttat ggaatacggt 60
 actggaacag ataaataatt tagttaataa catgacaaag aacagaaatt gtatacacta 120
 tacagcatag taatagaata atgaatgatt aaagttatta atattaggta gaaaatgaag 180
 ggtatctttg agagcagaac tcaaggaagc aagcaatttg ccttatgagg aaagagttac 240
 ctgtggataa aggagaaact gaaaaattta caagtcaaga ctttttgagc aaaaacaaaa 300
 atatgactat gagtcaccaa ttcagtacag tgaaaaaaaaa gttgaagaga tatcttggaa 360
 gtaaaccatg ttgtggaaga gcagggtttt gataatcatg ggattattct gaatgaattt 420
 taaatgcgat aggaatatat gagataattt caccagagaa taatatgatc atgtttgcat 480
 tt 482

<210> 603
 <211> 372
 <212> DNA
 <213> Homo sapien

<400> 603
 gttccaacct tcatttctga aactgttcta gagcactttg tctttctcgt agttcataac 60
 ttaccccttc agtctagaat tagaattaca ttatctgttt tactacttta ctgactgta 120
 agctcctaga agataaggac tagggagttc atctctgtat tccaccagaa ggtacagtga 180
 ctcataacta gagtcttttag atgaaactta ctgagttgaa taacttaata tatttctggt 240
 ttcatccca agggaggcca tgtctggaga tagaccttga atttaataaa ttttaggcac 300
 tataccattt cagtggagaa aattgttggg aaatttgggg ggatggatat ataaggggga 360

cctgctagct aaaaatatta tacccttgca aa

572

<210> 607
 <211> 178
 <212> DNA
 <213> Homo sapien

 <220>
 <221> misc_feature
 <222> (1)...(178)
 <223> n = A,T,C or G

<400> 607	
ctcggggtaa tctcccagca agaggtcagg tcctggntgt gcgtcccagg gtgtcagtga	60
aattggctgc tcccctgacc cagggcacct tcatgcgtct tcacagcagg actactgtga	120
ccaaggccag acctttcatc tttcaaaaga ctttgactaa aaatgcttta aaaaagca	178

<210> 608
 <211> 416
 <212> DNA
 <213> Homo sapien

<400> 608	
cctgtctttg aatggatgaa ataggttaat aaagaacatc actgttttaa aactagaaca	60
ctgaaaaatt ctaggaaagc ttattttccc ttatatTTTT atggtaacttt caacacttaa	120
taacactatt tcaattaagt tttctcctag agtttatagt atatcagtac attcctttct	180
gtggatgcaa taatatagaa tcttattcca aatcttactg gcaggttctc tttaaattctt	240
caacggctgt catagtgtatt aaccaaaatt agttatgatt tctgcctatc tgtgtgagaa	300
cttacagggg aaattgttct aaacctgagg aacatgaagt aactgtactg cacactccaa	360
atgatgacag tcattttata tcaccttcaa ttacccaaca gcttttaata gtctgg	416

<210> 609
 <211> 648
 <212> DNA
 <213> Homo sapien

<400> 609	
ctgatctctc agcagaaaact cttcaaacca gaagagagtg ggggcccaata ttcaacattc	60
ttaaagaaaa taattttcaa cccagaattt catatccagc caaactaacc ttcacaagtg	120
aaggagaaat aaaatccttt acagacaagc aaatgctgag agattttatc accaccaggc	180
ctaccctaaa agagttcctg aaggaagcac taaacatgga aaggaacaac cagtaccatc	240
gaggctagga agaaaccgca tcaactaagg agcaaaataa ccagctaaca tcataatgac	300
aggatcagat tcacacataa cgatattaac tttaaatgta aatggactaa atgctccaat	360
taaaagacac agactggcaa attggataaa gagtcaagac ccatcagggt gctgtattca	420
ggaaacccat ctcaccgtgc agagacacac ataggctcaa aataaagggc tggaggaaga	480
tctaccaagc aaatggaaaa caaaaaaagg caggggttgc aatcctagtc tctgataaaa	540
cagactttta accaacaag atcagaagag acaaagaagg ccattacata atggtaaaagg	600
gatcaattca acaagaagag ctaactatcc taaatatata ttgcaccc	648

<210> 610
 <211> 310
 <212> DNA
 <213> Homo sapien

```

<400> 610
ccagctcttc tctgtcacat tcctatttct gacttctgcc tggctttcag tttctgcccc      60
accttggttt tttcccagct tgaacctaat agaactccag agtttggggg gagggcccagc      120
cctttgtttt ctgctcttga agcatattca cacataaaaa gttgtattct cttacacaaa      180
ctgttttgag gctcttaccg tagtogaagg tatcttagat cttccttagt gatctcatta      240
agaatatccg aaagtgtata accctcttca acaatctgaa acaaagatca gatccttaag      300
agctgagcag                                     310

```

```

<210> 611
<211> 254
<212> DNA
<213> Homo sapien

```

```

<220>
<221> misc_feature
<222> (1)...(254)
<223> n = A,T,C or G

```

```

<400> 611
ctgtttttac atctaaagca atagactaga actgaattnt cttctacata gtaaaatcac      60
aattgtggaa ttacaggaat tctggtgata ttaagggtgaa acaacaaaac acaaaaggcc      120
ctattttaac agttgatgtg acagtaagtt ttaatagaac ctgtaacttc attttggaag      180
tgcttctcca ccaaataaag cctttttccc ctatttaagg agccagatgg attgaaagat      240
gtggaaatag gcag                                     254

```

```

<210> 612
<211> 225
<212> DNA
<213> Homo sapien

```

```

<220>
<221> misc_feature
<222> (1)...(225)
<223> n = A,T,C or G

```

```

<400> 612
ctgactatat catgtcacca tcatagccaa tacaacattn ttgccatact tcctaaaaac      60
cttttcgcat aactgatca tgctacttat cagcactttc taacatcctg accaaacaga      120
caccacacc tcttatagag tacactgtga gagaataaca tggacttgat atggcatcac      180
acttgtttta aagcaaaaaa aaaagaaaaa gaaaagaaaa aaaaaa                    225

```

```

<210> 613
<211> 471
<212> DNA
<213> Homo sapien

```

```

<220>
<221> misc_feature
<222> (1)...(471)
<223> n = A,T,C or G

```

```

<400> 613
ccatcagact tcttggtgct ctggctatat tcaatgtgaa gtaaaaaata tcccagtcct      60
tacacaaaaa tagaggctct gacttagaag tatgctttta gctttctttt taaataagac      120

```


tttttaccat tctaattggtt acaaagtaac cag

393

<210> 621
<211> 563
<212> DNA
<213> Homo sapien

<400> 621

ctgacaatga	taaaattatc	tctatatggg	caaacgcgtg	ctctttgtcg	aagaagaaag	60
cttcagcttc	atgttccagg	tgagttaatt	aggcaatgta	tgaatgctaa	tatctctttc	120
acataatttg	cttaagatct	gtcttaggac	tctcgtcttg	cccatatggt	tttccaaggg	180
cagaagggcc	tctttttgat	gagaggcagt	tttcagtaac	tcttaaagtg	ataacagcaa	240
aggagaggag	agagaagagt	aagacaaatc	gaaacattct	tcaattgctt	cttggccttt	300
tggctaagct	caagctcaaa	acaggtcttc	aaggagaaaa	tacatcacia	agaaaaggat	360
gttttatttc	ttaccttgct	ctagaaaaat	ttccataaac	tctattggct	taattctgta	420
aacttgacca	atatcagagt	gcttcctacc	aaggagggta	gctgatgagc	gtgaccatgg	480
tacatcctag	aagaatgtgt	gatgaagaag	ctttcacctg	gtaaaagagt	tgaaaattat	540
tcaaggagac	attatggtct	tgg				563

<210> 622
<211> 505
<212> DNA
<213> Homo sapien

<220>
<221> misc_feature
<222> (1)...(505)
<223> n = A,T,C or G

<400> 622

tcttaagtgt	gtttaataga	taaagtaaac	tttcctagtc	aagggttaga	tttttattat	60
ctcttggttt	ccgactttct	acttttcaac	tttgaacttc	aaaaaaacat	tactttgctt	120
atccttttga	ctttgatcag	gttggtttaga	attgtagatc	aaaccattct	ttgatcattt	180
tattgtttta	atgnttagtt	ccatttataa	tttttatagc	caactctcgg	ttattttctgt	240
cttttgagat	tgcaattcag	aagctgtatg	tcgaagtaat	ttatgagttg	actttttatac	300
ttaggcttct	ttaaatacta	atagtcaaga	attctagagc	atctaataaa	aaattaactt	360
tcagatcatt	gggaatctgt	cctcatttaa	atatgtgtaa	atgcatttcc	acagcaaatt	420
gcttcatgcc	ctttgnctat	aaggaaatta	ttccttgtag	ctaatacatt	tttcattttg	480
cagnccaaat	cttttttgag	aaagg				505

<210> 623
<211> 489
<212> DNA
<213> Homo sapien

<400> 623

cctactatgg	gtgttaaatt	ttttactctc	tctacaaggt	tttttcctag	tgtccaaaga	60
gctgttcctc	tttgactaa	cagttaaatt	tacaagggga	tttagagggg	tctgtgggca	120
aattttaaagt	tgaactaaga	ttctatcttg	gacaaccagc	tatcaccagg	ctcggtaggt	180
ttgtgcctc	tacctataaa	tcttccact	attttgctac	atagacgggt	gtgctctttt	240
agctgttctt	aggtagctcg	tctggtttcg	ggggtcttag	ctttggctct	ccttgcaaag	300
ttattttctag	ttaattcatt	atgcagaagg	tataggggtt	agtccttgct	atattatgct	360
tggttataat	ttttcatctt	tccttgccg	tactatatct	attgcgccag	gtttcaattt	420
ctatcgctat	actttatttg	ggtaaatgg	ttggctaagg	ttgtctggta	gtaagggtgga	480


```

ccatnngaac gcactcagga ggtgggttgt tctggatgca gaaaccagag atctagtttc      60
tatccacaca gacgggaatg aacagctctc tgtgatgcg tactcaatag atggtacctt      120
cctggctgta ggatctcatg acaactttat ttacctctat gtagtctctg aaaatggaag      180
aaaatatagc agatatggaa ggtgcactgg acattccagc tacatcacac accttgactg      240
gtcccagac aacaagtata taatgtctaa ctcgaggagac tatgaaatat tgtactggga      300
cattccaaat ggctgcaaac taatcaggaa tcgatcggat tgtaaggaca tttgattgga      360
ccgacatata cctgtgggct aggacttcca gga                                     393

```

```

<210> 628
<211> 233
<212> DNA
<213> Homo sapien

```

```

<220>
<221> misc_feature
<222> (1)...(233)
<223> n = A,T,C or G

```

```

<400> 628
ctggatttat aaaatagttg aatgacaaaa gaagnntggt ttgacagtaa aaaaaagaca      60
ttatggacaa aatatgcaaa atgtgcaaag aaaaaataaa tttgcattag aaaggtgggc      120
atttgatctc tgagccctgt gccatgtaac attgccatgt tctttcactg ttgtttgaat      180
gttgtagccc ancccttgac tctggactta aggcaagcta tgactggctt tgg                                     233

```

```

<210> 629
<211> 450
<212> DNA
<213> Homo sapien

```

```

<220>
<221> misc_feature
<222> (1)...(450)
<223> n = A,T,C or G

```

```

<400> 629
ccnggacaat ntaggcagga gaaggaaata aagggtattc aattaggaaa agaggaagtc      60
aaattgtccc tgtttgaga tgacatgatt gtatatctag aaaaccccat tgcctcagcc      120
caaaatctcc ttaagctgat aagcaactcc agcaaagtcg caggatacaa aatcaatgga      180
cacaaatcac aaacattctt atacaccaat aacagacaaa cagaggccaa atcacgagtn      240
gaactctatt ccaattgctt tcaagaaaat taaaatacct agggatccaa cttacaaggg      300
acatgaagga cctcttcaag gagaaactac aaaccactgc tcaatgaaat aaaagaggat      360
acaaagaaat ggaagaacat tccatgctca ttggtagctt gatgggggatg gcattgaatc      420
tataaattac cttgggcagt atggacctca                                     450

```

```

<210> 630
<211> 486
<212> DNA
<213> Homo sapien

```

```

<400> 630
cctactatgg gtgttaaatt ttttactctc tctacaaggt tttttcctag tgtccaaaga      60
gotgttcttc tttggactaa cagttaaatt tacaagggga ttttagagggt tctgtgggca      120
aattttaaagt tgaactaaga ttctatcttg gacaaccagc tatcaccagg ctcggtaggt      180
ttgtcgcttc tacctataaa tcttcccact attttgctac atagacgggt gtgctctttt      240

```



```

tttgttttac ataaaattag agtatgaaaa ccagtgttca attttttatac ttgttgagct 300
tgtaaaatgc cagcaattta aaactaggac ttttccccc ataagccaag gaggtagaat 360
tactaataca aggggttaaag aaggtagatt ttgttttcaa tatttgggta atattagaaa 420
gattcttccc acagggaaga actagcaagt gtcccaattt tttccaaacg ttggggaggg 480
gaaaattcac tgtatcatga aaccctaagg gtttgngtgc acttcctgct ttttagg 537

```

```

<210> 638
<211> 445
<212> DNA
<213> Homo sapien

```

```

<220>
<221> misc_feature
<222> (1)...(445)
<223> n = A,T,C or G

```

```

<400> 638
ccagcagaac acagnagtga tttgggtccc tttgttcccc agtgggggtat ctatccttgt 60
gcagggcaca agcctacatg gtggctctgg tcatatcatt agaaaataga cagaaatggg 120
ctgcacacca gaatgaatga attgaattga aaggaggagg tgatgggtgga aaaaaaaca 180
agtcaattca tttagactgg tagaaccaga accactgtgt agtacatcca aacgggttaa 240
attccctgga agatgttaca taatcctatc atgggtgttta tttatggaaa tctattttaa 300
aaattttatg taatactgca cagtctgttt gcatgatgcc ttgtacgtag tagcaactca 360
gtaaatactt tttgaatgaa ctagtatagt attttaatta gctagtcttc gtgtactgg 420
acaaaagaac agtgtcatct tacag 445

```

```

<210> 639
<211> 584
<212> DNA
<213> Homo sapien

```

```

<400> 639
gcttgagtat tctatagtgt cacctaaata gcttggcgta atcatgggtca tagctgtttc 60
ctgtgtgaaa ttgttatccg ctacacaattc cacacaacat acgagccgga agcataaagt 120
gtaaagcctg ggggtgcctaa tgagtgaagt aactcacatt aattgcgttg cgctcactgc 180
ccgctttcca gtcgggaaac ctgtcgtgcc agctgcatta atgaatcggc caacgcgcgg 240
ggagaggcgg tttgcgtatt gggcgctctt ccgcttcttc gctcactgac tcgctgcgct 300
cggtcgttcg gctgcggcga gcggtatcag ctactcaaaa ggccggttaata cggttatcca 360
cagaatcagg ggataacgca ggaaagaaca tgtgagcaaa aggccagcaa aaggccagga 420
accgtaaaaa ggccgcgttg ctggcggttt tccataggct ccgccccctt gacgagcatc 480
acaaaaatcg acgctcaagt caagagggtg cgaaaaccga caggactata aagataccag 540
gcgtttcccc ctggaagctc cctcgtgcgc tctcctgttc cgac 584

```

```

<210> 640
<211> 404
<212> DNA
<213> Homo sapien

```

```

<400> 640
ccataggaac gcactcaggc aggtgggttt ttctggatgc agaaaccaga gatctagttt 60
ctatccacac agacgggaat gaacagctct ctgtgatgcg ctactcaata gatggtaoct 120
tcctggctgt aggatctcat gacaacttta tttacctcta tgtagtctct gaaaatggaa 180
gaaaatatag gagatatgga aggtgcactg gacattccag ctacatcaca caccttgact 240
ggtccccaga caacaagtat ataatgtcta actcgggaga ctatgaaata ttgtactggg 300

```

acattccaaa tggctgcaaa ctaatcagga atcgatcgga ttgtaaggac attgattgga 360
cgacatatat ctgtgtgcta ggatttcaag tatttggtgt ctgg 404

<210> 641
<211> 138
<212> DNA
<213> Homo sapien

<220>
<221> misc_feature
<222> (1)...(138)
<223> n = A,T,C or G

<400> 641
ctgtgacagg aacattacct gaagtgcagg gtggttacct gcacaaagtc ccatttccaa 60
aaatttctgt gtaattcacc agaaattttg gatggaataa ttagaaaaaa aaaaagagggt 120
taaaacntgt aactcaaa 138

<210> 642
<211> 381
<212> DNA
<213> Homo sapien

<220>
<221> misc_feature
<222> (1)...(381)
<223> n = A,T,C or G

<400> 642
ctgtagggtg aatttttacc cagaaaagat aggccctaga agcctcattt cttttctcca 60
tggaaaagga cagccctctg ctgcagcgtt caacttgtgt gtttactgac agagtgaact 120
acagaaatag cttttcttcc taaaggggat tgttctacat tttgaagtta ttttttaata 180
aaattgaatt atgttgtgta ttgtgcttcc taataggaaa tgcattattg gactgttttt 240
gtaacatcct gtttattgca aatagctagt atcgttcaaa aactgtataa aatacttttg 300
tacatattag caatgtctaa tttgtataca cttcagttaa atttccttaa aacttgaaag 360
gggaccttgt anaaattaaa a 381

<210> 643
<211> 403
<212> DNA
<213> Homo sapien

<400> 643
ccttcctaaa aaatagtggg gagctggagg ctacttccgc cttcttagcg tctgggcaga 60
gagctgatgg atatcccatt tggccccgac aagatgacat agatttgcaa aaagatgatg 120
aggataccag agaggcattg gtcaaaaaat ttggtgctca gaatgtagct cggaggattg 180
aatttcgaaa gaaataattg gcaagataat gagaaaaagaa aaaagtcatt gtaggtgagg 240
tggttaaaaa aaattgtgac caatgaactt tagagagttc ttgcattgga actggcactt 300
attttctgac catcgctgct gttgctctgt gagtcctaga tttttgtagc caagcagagt 360
tgtagagggg gataaaaaga aaagaaattg gatgtattta cag 403

<210> 644
<211> 688
<212> DNA


```

aaaaaaaaagg cgaactctgc cttggagagg tagatgataa gaaataaaaa ggtgtttata 120
actattttgt attataaagt gggccttaga gataggaaga agaatzgatgg attccttttg 180
gatcaatcag aaaggaaaca cgaaagaaaa gtcaggaagg tagagagaga aaaagggagg 240
gaaggagaaa gaatgggaat aaaataagga ggtaagagat actatttttg ctgagcaacc 300
agtgtgtttc aggatgatac aaagaaaaat atagaataga aataagtgca ggcttggaat 360
cagctacaaa tcctaaagat ggggtgtgtg tggatgtgtg tgtgtgtgtg tgnacaccat 420
tgtgtgtttg taaaatgtgt atgtccc 447

```

```

<210> 647
<211> 388
<212> DNA
<213> Homo sapien

```

```

<400> 647
gaaggtgata taaaatgact gtcattcattt ggagtgtgca gtacagttac ttcattgttcc 60
tcaggttttag aacaattttcc cctgcaagtt ctccacacaga taggcagaaa tcataactaa 120
ttttggttaa tcactatggc agccgttgaa gaatttaaga gaacctgcca gtaagatttg 180
gaataagatt ctatattatt gcatccacag aaaagaatgt actgatatac tataaactct 240
aggagaaaac ttaattgaaa tagtgttatt aagtgttgaa agtaccataa aaatataagg 300
gaaaataagc tttcctagaa tttttcagtg ttctagtttt taaacagtga tgttttttat 360
taacctattt catccattca aagacagg 388

```

```

<210> 648
<211> 632
<212> DNA
<213> Homo sapien

```

```

<220>
<221> misc_feature
<222> (1)...(632)
<223> n = A,T,C or G

```

```

<400> 648
cctggctggg cntttgacct gcgnttttaa atnactcaca gaggggtggga caggaggaag 60
agtgaaggaa aaggtcaaac ctgttttaag ggcaacctgc ctttgttctg aattggtctt 120
aagaacatta ccagctccag gtttaaattg ttcagtttca tgcagttcca atagctgac 180
attgttgaga tgaggacaaa atcctttgtc ctactagtt tgctttacat ttttgaaaag 240
tattattttt gtccaagtgc ttatcaacta aaccttgtgt taggtaagaa tggaatttat 300
taagtgaatc agtgtgacct ttcttgatc aagattatct taaagctgaa gccaaaatat 360
gcttcaaaag aagaggactt tattgttcat tgtagtcat acattcaaag catctgaact 420
gtagtttcta tagcaagcca attacatcca taagtggaga aggaaataga tagatgtcaa 480
agnatgattg gtggagggag caaggttgaa gataatctgg ggttgaaatt ttctagttnt 540
cattccgtac attttttagtt agacatcaga tttgaaatat taatgttacc tcctcaatgg 600
ggtggtatca gacctgcccg ggcgncgcn tc 632

```

```

<210> 649
<211> 300
<212> DNA
<213> Homo sapien

```

```

<220>
<221> misc_feature
<222> (1)...(300)
<223> n = A,T,C or G

```

```

<400> 649
nggtgaagat agaanaaata taagcgaaat tggataaaat agcactgaaa aaatgaggaa      60
attatttggt accaatttat tttaaaagcc catcaattta atttctggtg gtgcagaagt      120
tagaaggtaa agcttgagaa gatgaggggtg tttacgtaga ccagaaccaa tttagaagaa      180
tacttgaagc tagaagggga agttgggttaa aaatcacatc aaaaagctac taaaaggact      240
ggtgtaattt aaaaaaaaaact aaggcagaag gctttggaag agttagaaga atttggaagg      300

```

```

<210> 650
<211> 498
<212> DNA
<213> Homo sapien

```

```

<220>
<221> misc_feature
<222> (1)...(498)
<223> n = A,T,C or G

```

```

<400> 650
ngtntctgnta aacagaaggg tacaangccc ttctggcttt aagcagtcac aggaatgtga      60
cagacattcc tcttagggag cgcctcctcc taggggttcc tcatctgtct cacactgagt      120
ggatgtaatg ctattttaat cctgctgtgg cccccaatac tagtacttgt ccataccttc      180
ttgcattttt agcgtctgct ctgtgggggtt gttaggccct ggcactccca ggaactagtg      240
ctaaagctgc atctntctct cccctctagg gatcgataaa gtttccactgc agaaagtctc      300
cactgcggta tgctgacatc tgcctgaac cttcacccta cagcattaca ggctttaatc      360
agattctgct ggaaagacac aggetgatcc acgtgacctc ttctgccttc actgggctgg      420
ggtgacccct ggtgcctttg tttccacaag gccttttcct gcccctgccc ttgccaaaga      480
catttaataca gcacacag

```

```

<210> 651
<211> 654
<212> DNA
<213> Homo sapien

```

```

<220>
<221> misc_feature
<222> (1)...(654)
<223> n = A,T,C or G

```

```

<400> 651
ctgaggggtcc ccagggtttct aaagctctca ggacgagaaa gtaggtccca agataaggag      60
cctaaagggc ttttttcttt ctgtgtattc cttcttggcc tccaacatgg gtacagtac      120
aagagcatgt aacagagaag aaggactana cctaccattt tctggataaa gaattggaaa      180
gaggatccac aggtaaccaa aaagtaccag ggaaatggca gagaaggaaa acctcaggag      240
accaacctca taagtgggat ttattagncc ctgggctcaa atccaaattg tacatgaata      300
tgtctgggtcc tagatagggt accgaagact ttgaaagtga attttggtat atcattgccc      360
agattccaga ctggnatttg tgtgacacaa catacaggat atatctgaat agtgctcaga      420
agagtttgaa aatgcaaattg atattaaaat aaagatgaaa aagagaaaagc tggtcagaac      480
ttgtggacat aacccttctg gatctgtngc ctgattaaaa aatagttgat attctcgaat      540
gaattaaaac aagatttaga gactgagcat ggtagctnat tcttgtaatc caacnctttg      600
ggaggggcaag gcaanagaat tgcttgccgc caggagtttt gagaccagct tggg      654

```

```

<210> 652
<211> 293

```

<212> DNA
<213> Homo sapien

<220>
<221> misc_feature
<222> (1)...(293)
<223> n = A,T,C or G

<400> 652
ngtctgttgc actgaggtga ctaaggatac attttgagga agtagctcca agaacatttc 60
cattttcact gtgccttcac atacatctaa tggaaatgaa cagcaccctt catccatcca 120
cggaagcgat taagaaaagg gtgggatgga aaaattaacc caacaatatt agatcaatac 180
gtagtattta agngtccata atgtgccagg ctgaagatgc acgggaaaac cacactagcc 240
ggtctgtcaa gggcttgaga ataccataaa caagaaaaca gacgaaccaa ttt 293

<210> 653
<211> 294
<212> DNA
<213> Homo sapien

<220>
<221> misc_feature
<222> (1)...(294)
<223> n = A,T,C or G

<400> 653
ngtccaccac tgcagcccta catacagttg aaaaaaaatt ccattctggtt aacatttggtt 60
ttataagttt tcacgcaata cacaaaaaac ccctctgcac ttcttgtaaa gaacaaaaaa 120
gatacacaa agttaagcgt aaagatcaca ggcaatagca ttcaaactatg gatgtgggta 180
gagaaaggag tacctggcat gagtacctgc ttagtttgac tgaatccttg atttttaatt 240
tggcttttca tggggccgctc acaacaccaa cgctgtgtga ggtatggtag tcag 294

<210> 654
<211> 250
<212> DNA
<213> Homo sapien

<400> 654
ctgtccttga acaagtatca atgtgtttat gaaaggaaga tctaaatcag acaggagttg 60
gtctacatag tagtaatcca ttgttggaat ggaacccttg ctatagtagt gacaaagtga 120
aaggaaattht aggaggcata ggccatttca ggcagcataa gtaatctcct gtcctttggc 180
agaagctcct ttagattggg atagattcca aataaagaat ctagaaatag gagaagattt 240
aattatgagg 250

<210> 655
<211> 494
<212> DNA
<213> Homo sapien

<400> 655
ccattataat tttataaac cattaccctt taaattctac cgattataag cagcgtaaaa 60
gtaactatat aaagcaaaca tcgcaaagga actctgcagg agctcttaatt tcctttatgt 120
agctatcata aaattcactt tcctgaagac atttactctc attcacttcc aaactccaaa 180
cctttttctg gtagcaccac ttttgttttt aatagaaaga tgagttcata tctgtacatc 240


```

cctgaaaaga aagntgctct tatggactct tgcattgttaa gactatgtct tcacatcatg      60
gtgcaaataca catgtaccca atgactccgg ctttgacaca acaccttacc atcatcatgc      120
catgatggct tccacaaagc attaaacctg gtaaccagag attactggtg gctccagcgt      180
tgtagatgt tcatgaaatg tgaccacctc tcaatcacct ttgagggcta aagagtagca      240
catcaaaagg actccaaaat cccataccca actcttaaga gatttgtcct ggtacttcag      300
aaagaatttt catgagtgtt ctttaattggc tggaaaagca ccag                          344

```

```

<210> 659
<211> 230
<212> DNA
<213> Homo sapien

```

```

<400> 659
ctgctttccc tgctaaacag ttccagagca aaagcagcaa aaagaaaata tgggagggat      60
atgggcaacg tatactcgaa cgtacgcaga gaagagagta cggtttagctc taatatttct      120
cattgaactt ggtggtatgt gccttccctg catataaggc catagtgtctt ttttgggagc      180
gctagaatat ccatccactt gacagtgtacc acaaaatagg ctgtttccag                230

```

```

<210> 660
<211> 80
<212> DNA
<213> Homo sapien

```

```

<400> 660
ctggtccttg ttaaactcga tcaccacttt ggagagatcg actggagggt cctgggtgtt      60
ctgagggggc tgggggacag                          80

```

```

<210> 661
<211> 535
<212> DNA
<213> Homo sapien

```

```

<220>
<221> misc_feature
<222> (1) ... (535)
<223> n = A,T,C or G

```

```

<400> 661
ctgaaccata tctgattaac tctttggtct ctgttatttg aacaaaaccg acgctatgcc      60
tgcagccgcc agactgcaac caaaaacaca gtttggggtc agaagacatt aaaaatcaca      120
ataaaatagg atgaatgttc taagtcacgc aactgaatca aggcaccttt ttttttcaaa      180
agcaaaaagt tgtttaacaa tattccagaa tagtagatac ttcaaaaacc agattacagt      240
atatatcatt ttgctgcaca ttttagtcta ttttctgtat acatagtcac acattcttta      300
ccctctccca acttatacat gctttatccc cccagtcacg tgctatgtag gtataaaaaa      360
ataaagtgtg atctaaacaa gtgattttaa aaaaaaaact aacgaatgcc ncnatnataa      420
cnctgaactt gtttccctnt tgaaggacat tggaaatgtt accgaggtn ntttacctng      480
gccgcaaccn cnctangggc naattccagc ncactggggg ccgttactag gggat                535

```

```

<210> 662
<211> 257
<212> DNA
<213> Homo sapien

```

```

<400> 662

```

```

cctgactaaa gcacatatca cactccctac acttccatgt tttctctccc atgtggaccc      60
tctgatgcat atcaagattc aagcgctgt tgtagccctt cccacagtcc tcacatttgt      120
atgggttttc tacactgtga actttttctt gcactttaga gaatgaattc tgtacaatgt      180
tcttcccatg ctgctcacat ttgagagggtg tttctctgct gtggcgtctc tgatgggtca      240
gacgagttga ggaccag                                     257

```

```

<210> 663
<211> 516
<212> DNA
<213> Homo sapien

<220>
<221> misc_feature
<222> (1)...(516)
<223> n = A,T,C or G

```

```

<400> 663
ccaattatag gtatttttatt ttttaaagat tagagngttc ttgaagctct ttctatttct      60
ttgtcaatga actaaacatt ggcaaatatg tagggtttcc cacataagaa cattattaac      120
atcaaaatag aaagctggtg gtgaaaataa tgattgggaa cacagagtct ctactcagcg      180
ttctacttct gccataccat aactttgtga tctcacgaaa tatctctcca tgttctcatc      240
cctatgtata gttctgtcat ttttcaataa gagctttttg ctttaattatg aagtactagt      300
tactataacc attattttga gcttcatgta aatcaagaac acatggactc cacttgcaaa      360
acattgaaaa tgtagttagg gattgggggc aaaaagcaac atttttaaatt gtgtaaagac      420
aatgagtaag caacaaagtg tccaattttt taggcgaaag ttgcatatgt caggaaaagg      480
caggattaag taatagagaa tttgaatgat aactgg                                     516

```

```

<210> 664
<211> 212
<212> DNA
<213> Homo sapien

```

```

<400> 664
gtccgaggag gttagtgtgt gcaataaaaa tgattaagga tactagtata agagatcagg      60
ttcgctcttt agtgttgtgt atggctatca tttgttttga ggtagtttg attagtcatt      120
gttgggtggt aattagtcgg ttgttgatga gatatttgga ggtggggatc aatagagggg      180
gaaatagaat gatcagtact gcggcgggta gg                                     212

```

```

<210> 665
<211> 408
<212> DNA
<213> Homo sapien

```

```

<220>
<221> misc_feature
<222> (1)...(408)
<223> n = A,T,C or G

```

```

<400> 665
atccaggggt ncccggtngc tgcngggaaa cctccagcct tgttcttcaa accactcagc      60
tcatgtgttt tgcgctgact agtactgaat aatacaacca ctcttattta atgttagtat      120
tatttttttg acaactcagt gtctaacagc ttgatatgca ggtccttgca tcttacattt      180
cttttaggaag ttaccatttt gtaactttta aaacaggaaa aatatcagtt ggcaaatgca      240
atcttttttt tttttaagct aaaggggggn naacngnaan naaaatnttt ntgangtnng      300

```

gtctataagc acccttggang ggatntgtta aaagnngncat naanggggga ttctcntttt 360
gcaaaaaaat ntaannatca atttatanan ctttatTTTT nactttnt 408

<210> 666
<211> 635
<212> DNA
<213> Homo sapien

<220>
<221> misc_feature
<222> (1)...(635)
<223> n = A,T,C or G

<400> 666
ctgaagnaca agggtcaggc aaaaataaga tcacaatcac caatgaccag aatcgcttga 60
cacctgaaga aatcgaaagg atgggttaatg atgctgagaa gtttgctgag gaagacaaaa 120
agctcaagga gcgcattgat actagaaatg agttggaaag ctatgcctat tctctaaaga 180
atcagatttg agataaagaa aagctgggag gttaaaccctc ctctgaagat aaggagacca 240
tggaaaaaag tgtagaagaa aagattgaat ggctggaaag ccaccaagat gctgacattg 300
aagacttcaa agctaagaag aaggaaactgg aagaaattgt tcaaccaatt atcagcaaac 360
tctatggaag tgcaggccct cccccaactg gtgaagagga tacagcagaa aaagatgagt 420
tgtagacact gatctgctag tgctgtaata ttgtaaatac tggactcagg aacttttggt 480
aggaaaaaat tgaaagaact tanctctcga atgtcattgg aatcttcacc tcacagtggg 540
gttgaaactg ctatagccta agcnggctgt ttactgnttt ncattagcag gtgctcacca 600
tgtctttggg gtgggngggg ggagaaagaa agaan 635

<210> 667
<211> 388
<212> DNA
<213> Homo sapien

<400> 667
gaagggtgata taaaatgact gtcattcattt ggagtgtgca gtacagttac ttcattgttcc 60
tcagggttag aacaatttcc cctgtaagtt ctacacagaa taggcagaaa tcataactaa 120
ttttgggttaa tcaactatggc agccgttgaa gaatttaaga gaacctgcca gtaagatttg 180
gaataagatt ctatattatt gcatccacag aaaagaatgt actgatatac tataaactct 240
aggagaaaaac ttaattgaaa tagtgttatt aagtgttgaa agtaccataa aaatataagg 300
gaaaataagc tttcctagaa tttttcagtg ttctagtttt taaacagtga tgttttttat 360
taacctattt catccattca aagacagg 388

<210> 668
<211> 498
<212> DNA
<213> Homo sapien

<220>
<221> misc_feature
<222> (1)...(498)
<223> n = A,T,C or G

<400> 668
tgatcttaac aaaattcgta gcagtggaaac cttgaaatgc atgtggctag atttatgcta 60
aaatgattct cagtttagcat tttagtaaca cttcaaagggt ttttttttgt ttgttttcta 120
gacttaataa aagcttagga ttaattagaa gaagcaatct agttaaattt cccatttgta 180

ttttattttc	ttgaataactt	ttttcatagt	tattcgttta	aaaagattta	aaaatcattg	240
cactttggtc	agaaaaataa	taaatatatc	ttatgaatgt	ttgattccct	tccttgctat	300
ttttattcag	tagatttttg	tttggcatca	tgttgaagca	ccgaaagata	aatgattttt	360
aaaaggctat	agagtccaaa	ggaatgttct	tttacaccaa	ttcttccttt	aaaaatntct	420
gaggaatttg	ttttcgctt	actttttttt	cttctgtcac	aatgctaagn	ggtatccgag	480
gtntttaata	tgagattt					498

<210> 669

<211> 622

<212> DNA

<213> Homo sapien

<400> 669

ccttagccaa	agaatgcagt	ggagccttcc	cccttcaact	gcattgtgaa	tgaataccaa	60
ttaacagcat	aaaaattaat	agtcccatat	cagatctgga	aggggtttct	ggggctgtct	120
gatgtcccta	tcctgttgta	gtgaacacaa	tagcagaaaa	ttctttctgg	gtccatctgc	180
tataaagtct	tggtaaaaca	gcattactat	gaagaggatg	aactcaccta	ccttcagatg	240
gaggaaaagt	gaaaaggact	taggctttag	tcctccatga	cttttcttaa	gcactaccta	300
cctgtaataa	gctgagtgc	aaaggatgcc	gaagaaaatc	tgacccaga	agctgttaga	360
aagcactgca	gagaacaggg	tatgaagaaa	ataaagagtt	cttaataaac	ccttaagatt	420
ctttgttcaa	ggtaaccttg	ccaaaagggc	agagtaggtg	gcaaagagtt	gcttttaatc	480
tagctctaca	ctgcatttga	aaataaaatt	tgcccatttt	gaatatattg	tttataatta	540
aatgtgcttt	ttacactgca	ggtcaatata	aaaactgggt	agtaaatttc	cagcgagcat	600
ttatgttcat	ttgctcacag	ca				622

<210> 670

<211> 477

<212> DNA

<213> Homo sapien

<400> 670

ttgggccctc	tagatgcagt	ctcgagcggc	cgccagtgtg	atggatatct	gcagaattcg	60
cccttgccgc	ccgggcaggt	gatggatgag	gagcaaaaac	tttatacggg	tgatgaagat	120
gatatctaca	aggctaataa	cattgcctat	gaagatgtgg	tcggggggaga	agactggaac	180
ccagtagagg	agaaaataga	gagtcaaacc	caggaagagg	tgagagacag	caaagagaat	240
atagaaaaaa	atgaacaaat	caacgatgag	atgaaacgct	cagggcagct	tggcatccag	300
gaagaagatc	ttcggaaaga	gagtaaagac	caactctcag	atgatgtctc	caaagtaatt	360
gcctatttga	aaaggttagt	aaatgctgca	ggaagtggga	ggttacagaa	tgggcaaaat	420
ggggaaaggg	ccaccaggct	ttttgagaaa	cctcttgatt	ctcagtctat	ttatcag	477

<210> 671

<211> 127

<212> DNA

<213> Homo sapien

<400> 671

gtgtgtgtgt	ctacttgggc	gtgtttaacg	tgtgcgtttg	tgtctgcgtg	tgcatgtgtc	60
tgtgtgtgcg	cgtgtatttc	agtttggtt	gccggatccc	atatgattgc	gtgcctgtgt	120
acctgag						127

<210> 672

<211> 400

<212> DNA

<213> Homo sapien

<400> 672
 gggctctgcac agctatgtta acagcatcct tataccagga gtaggaggaa agacacgact 60
 ggaaaagcaa ttcaagctgg tcacacagtg taatgcaaaa tatgtggaat gtttcagtgc 120
 tcagaaagag tgtaacaaag aaaagaacag aaactcttca gttgtgccat ctgagcgtgc 180
 tcgagtgggt cttgcacat tgcttggaaat gaaaggaaca gattacatta atgcttctta 240
 tatcatgggc tattatagga gcaatgaatt tattataact cagcatcctc tgccacatac 300
 tacgaaagat ttctggcgaa tgatttggga tcataacgca cagatcattg tcatgctgcc 360
 agacaaccag agcttggcag aagatgagtt tgtgtactgg 400

<210> 673
 <211> 600
 <212> DNA
 <213> Homo sapien

 <220>
 <221> misc_feature
 <222> (1)...(600)
 <223> n = A,T,C or G

<400> 673
 ctggcggttg tcattagtga atgtatgaca gcaggatgtg aggggatgcc caggagtcag 60
 tgtagcatt gtcactgag atcactgcta ttaatatcat ccattaattt attagtgagc 120
 ttcactatat gcagactggg agataaggag aaaatctgtc acattctctc tagctaatac 180
 gatcagctac caattaatga gattctgaat gaaatatcaa tatgtgtttt tctaatttgg 240
 acctaggaca gagctgttg tttgtcataga gaaaaacaat aatgcttaaa catagcacat 300
 tataattaaa gcaggtttct cacatacttt tcattttatc ctttggataa ttttgtgagg 360
 aacgcaggac accaacttcc ctttcataga tacaatcccc atgctattga tgaaagtgtt 420
 tttgaatgaa gccatacaac aaataactga tcaaagtggc attacacca aatttcttag 480
 taggactcct gcatagaatg tttagataga cgtgaaaagt ttgttcanga ggaccagcaa 540
 gagagaaact gggttctttg ggagggtttc ggtgctacat ttataccctn catcagagtn 600

<210> 674
 <211> 140
 <212> DNA
 <213> Homo sapien

<400> 674
 ggtgggttgg gtaaagtgt gaggcaggag tccgaggagg ttagttgttg caataaaaat 60
 gattaaggat actagtataa gagatcaggt tcgtccttta gtgttggtga tggctatcat 120
 ttgttttgag gttagtttga 140

<210> 675
 <211> 245
 <212> DNA
 <213> Homo sapien

<400> 675
 gttgggttgg tgggtgtaaat gagtgaggca ggagtcaggag gaggttagtt gtggcaataa 60
 aaatgattaa ggatactagt ataagagatc aggttcgtcc ttagtggttg tgtatggcta 120
 tcatttgttt tgaggttagt ttgattagtc attgttgggt ggtaattagt cggttgttga 180
 tgagatattt ggaggtgggg atcaatagag ggggaaatag aatgatcagt actgcggcgg 240
 gtagg 245

<210> 676
 <211> 621
 <212> DNA
 <213> Homo sapien

 <220>
 <221> misc_feature
 <222> (1)...(621)
 <223> n = A,T,C or G

<400> 676
 ctgtccccag ggnaaatagt ngaattcaac taagatctgt taataagatg tcagaataac 60
 taataatttt attaggaaaa aatcatgttt taaatttcaa aatgacactt atttgtcaag 120
 taatatgatac ttggaaaatt ttaaagaaaa ataatacctac ttataaacta cttttttata 180
 attgttttca gaaaaaaagt ttacagtctt aaggaaaata ttcagggtcta tcatatgggt 240
 tgacagattt tttaaaagt atttttggtt aggtcttctt ttagaaaaaa attaacttca 300
 aggggttttt gtaccactat aatctctaata acttactcag aattactgtg tatttactta 360
 atttcttatt atgtgcctta ttatgtgctt aagatacaat aggttagagt ttaatctaaa 420
 tatcttgaaa gctatattgt gggcttggtt agcattttgt tttttctttc tctgttttgg 480
 taaggattta aaattttttt cattgcaatt ttaagtgggt ttcaataagt aatagttttt 540
 atcaaatttt tgggtgcttgg tgcagagacg gcgtggggaa ggggtgaatgg ttttgggaat 600
 aattcagtgc acacctgggg g 621

<210> 677
 <211> 210
 <212> DNA
 <213> Homo sapien

 <220>
 <221> misc_feature
 <222> (1)...(210)
 <223> n = A,T,C or G

<400> 677
 tttacataaan atattatcag catttaccat ctcacttcta ggaataactag tatatcgtct 60
 acacctcata tcttccctac tatgcctaga aggaataata ctatcactgt tcattatagc 120
 tactctcata accctcaaca cccactccct cttagccaat atttgtccta ttgccatact 180
 agtcttttggc gcttgccaag cagcggttagg 210

<210> 678
 <211> 383
 <212> DNA
 <213> Homo sapien

 <220>
 <221> misc_feature
 <222> (1)...(383)
 <223> n = A,T,C or G

<400> 678
 gtaggagtca ggtagttagg gttaacgagg gtggttaagga tggggggaat tagggaagtc 60
 aggggttaggg tgggttatagt agtgtncatg gttatttagga aaatgagtag atatttgann 120
 aactgattaa tgtttggggnn tgagttnta tatcacagcc anaattntat gatgnaccat 180
 gtanogaaca atgctacagg gatgaatatt atggagaagt antctanttt gaagcttagg 240

gagagctggg ttgtttgggt tgnnggetcan tgtcagttcc anataataac ttcttgggtct 300
 aggcacatga atattgttgt ggggaanaga ctgataataa aggtggatgc gacaatggat 360
 ttacataat gggggtatna gtt 383

<210> 679
 <211> 371
 <212> DNA
 <213> Homo sapien

<400> 679
 aaaatgaaaa tattgacaag agtttcagat agaaaaatgaa aaacaagcta agacaagtat 60
 tggagaagta tagaagatag aaaaatataa agccaaaaat tggataaaat agcactgaaa 120
 aaatgaggaa attatttgta accaatttat tttaaaagcc catcaattta atttctgggtg 180
 gtgcagaagt tagaaggtaa agcttgagaa gatgaggggtg ttacgtaga ccagaaccaa 240
 tttagaagaa tacttgaagc tagaagggga agttgggttaa aaatcacatc aaaaagctac 300
 taaaaggact ggtgtaatat aaaaaaaact aaggcagaag gcttttggaa gagttagaag 360
 aatttgggaag g 371

<210> 680
 <211> 176
 <212> DNA
 <213> Homo sapien

<400> 680
 cctaggattg tgggggcaat gaatgaagcg aacagatttt cgttcatttt gggttctcagg 60
 gtttggttata attttttatt tttatgggct ttggtgaggg aggtaagtgg tagtttgtgt 120
 ttaatatatt tagttgggtg atgaggaata gtgtaaggag tatgggggta attatg 176

<210> 681
 <211> 152
 <212> DNA
 <213> Homo sapien

<400> 681
 ctggagatgg atatgagact agtcaagatg tgaatgctaa ttggagagaa atataatttt 60
 aggaagatgc acattgatgt ggggttttga tgtgtctgat ttgactact caagctctgt 120
 ttacagaaga aaattgaatg gcgaggggtg gg 152

<210> 682
 <211> 141
 <212> DNA
 <213> Homo sapien

<400> 682
 ccagtgcctt cttgccgtgg tttagtgtt ggggtgttaga aataaaaaact caggtctatt 60
 tcttaccagt cagtaacaat ttttagagaa tgtacttggt atataatata tggacttcag 120
 gaactttgtt ggggtggggg g 141

<210> 683
 <211> 308
 <212> DNA
 <213> Homo sapien

<400> 683

```
<210> 684
<211> 277
<212> DNA
<213> Homo sapien
```

```
<210> 685
<211> 457
<212> DNA
<213> Homo sapien
```

```
<220>
<221> misc_feature
<222> (1)...(457)
<223> n = A,T,C or G
```

```
<210> 686
<211> 234
<212> DNA
<213> Homo sapien
```

```
<220>
<221> misc_feature
<222> (1)...(234)
<223> n = A,T,C or G
```

<400> 686						
ntggattttat	aaaatagttg	caatgacaaa	agaagtatgt	tttgacagta	aaaaaaagac	60
attattggaca	aaatatgcaa	aatgtgcaaa	gaaaaaataa	atttgcatta	gaaagggtggg	120
catattgatct	ctgagccctg	tgccattgtaa	cattggcaatg	ttctttcact	gttggtttgaa	180
tqttgtaccg	cagcccttga	ctctggactt	aaggcaagct	atgactggct	ttgg	234

<210> 687
 <211> 315
 <212> DNA
 <213> Homo sapien

<220>
 <221> misc_feature
 <222> (1)...(315)
 <223> n = A,T,C or G

<400> 687
 nngtctgtga aaaactcttt ggatgattct gccaaaaagg tacttctgga aaaatacaaa 60
 tatgtggaga attttgggtct aattgatggg cgctcacca tctgtacaat ctctgtttc 120
 ttgccatag tggctttgat ttgggattat atgcaccct ttccagagtc caaaccggt 180
 ttggctttgn gtgtcatatc ctattttgtg atgatgggga ttctgacct ttataacctca 240
 tataaggaga agagcatctt tctcgtggcc cacaggaaag atcctacagg aatggatcct 300
 gatgatattt ggag 315

<210> 688
 <211> 522
 <212> DNA
 <213> Homo sapien

<220>
 <221> misc_feature
 <222> (1)...(522)
 <223> n = A,T,C or G

<400> 688
 ctgaattaga ggaggagaaa agaagccatt nnggagtact ttaattgttt agatgtgaga 60
 ggctgaatgt ttgggttaag atgttagttg tcagaatcat gagaaaagg tttaagcaag 120
 gggcatttct aattctaaaa ataacaacta ctgttattta ttgagcacta tctttttgtt 180
 gggactgtgc taaagtactt gatttatattt ttaaaacctt acaaaaaact tacaaggtag 240
 gtactgaaag attcagtaat ttgttcaaag tcacacagca aataagcaac agactctgga 300
 tttgaaccag gcaatcctag agcctgtact gttagtaatt atacttttagc acctgtcaag 360
 aattcctgtt gagtgtcaag aagcaancac caagttagga tttaaagcaa acatgattga 420
 agaatactgt ggtgtggttg acagtagtgc ctaagtctgt tttcagagtg aaaaatgaca 480
 aattagattt taagtatggt ttggagataa tatcaggaca gt 522

<210> 689
 <211> 158
 <212> DNA
 <213> Homo sapien

<220>
 <221> misc_feature
 <222> (1)...(158)
 <223> n = A,T,C or G

<400> 689
 tctcaactta ntntnatacc cacaccacc caanaacagg gtttgtagg nattgtttgc 60
 attaataaat taaagctcca tagggctctt tcgtcttgc gtgtcatgcc cgctcttca 120
 cgggcagggtc aatttcactg gttaaaagta agagacag 158

<210> 690
 <211> 300
 <212> DNA
 <213> Homo sapien

<220>
 <221> misc_feature
 <222> (1)...(300)
 <223> n = A,T,C or G

<400> 690
 tagaactcgt atttttaaac ttctattctc tanccttttc cactacatta tgacacaaga 60
 ccttgcagaa agtcgtctgg aaaatatcag accatctctt acttgtccca tccaatctta 120
 catcgaatta tatgcaccct taaaaagtta tttggagttt taaaaaactc tattagccca 180
 aattacctga aataaaactcc tggcttggtc ccctaagtgt tataaaaaat tgattgaaaa 240
 tattcatttt aaaaatgaag ntcttgaatt tattttaaatt actgtcttgc agtgagttgg 300

<210> 691
 <211> 305
 <212> DNA
 <213> Homo sapien

<400> 691
 ctgttcagaa agctcattgg acctggtttt gaaaataaaa caaagttaaa accctggggag 60
 gagttattgt gcagtgtgga gtactcaggc tttcttataa agaaaaaaaa agttatctgg 120
 taccaaagtg tgcaacctac agaccctcag gtactgccct gtgacttctc tgtatgacat 180
 cacaaggctg ccaagtgcct gtttttctag aactaggagt tggtagaggt ttggtagtgc 240
 tgaaaccatg cataggattg gtttactaaa ttaaaacctt attacgtacg tcctccaaaa 300
 gacag 305

<210> 692
 <211> 582
 <212> DNA
 <213> Homo sapien

<400> 692
 caggaaatgg ataaccattt taactgtatt ttttgcagcc cgtaccttct tgggaataca 60
 attgtctaac tttttatttt tgggtctggc gttgtggtgt gcaaaactcc gtacattgct 120
 attttggcac actgcaacac cttacagatg tggaagatgt gaaatttgtc atcaattatg 180
 actaccctaa ctctcagag gattatattc atcgaattgg aagaactgct cgcagtacca 240
 aaacaggcac agcatacact ttctttacac ctaataacat aaagcagggtg agcgacctta 300
 tctctgtgct tcgtgaagct aatcaagcaa ttaatcccaa gttgcttcag ttggtcgaag 360
 acagaggtgc aggttaaggat gactgatagg aaatgttggg agttacgagt cacatcgttg 420
 tctacaaatc cattttaaag gtattggagg gtgagtaaaa ccttgaatgt gaaaaactta 480
 gctgaaaaat tgtaaaaaaca tttcacgcct accatgaata gatctgtttc tttctgtcca 540
 caatgatttg tgtcatagac ataattgatc aatttgcaat tg 582

<210> 693
 <211> 275
 <212> DNA
 <213> Homo sapien

<400> 693

```

ccaattgatt tgatggtaag ggaggggatcg ttgacctcgt ctgttatgta aaggatgcgt      60
agggatggga gggcgatgag gactaggatg atggcgggca ggatagttca gacggtttct      120
atttcctgag cgtctgagat gttagtatta gttagttttg ttgtgagtgt taggaaaagg      180
gcatacagga ctaggaagca gataaggaaa atgactatga gggcgtgatc atgaaagggtg      240
ataagctctt ctatgatagg ggaagtagcg tcttg                                     275

```

```

<210> 694
<211> 397
<212> DNA
<213> Homo sapien

```

```

<220>
<221> misc_feature
<222> (1)...(397)
<223> n = A,T,C or G

```

```

<400> 694
nggtctgcat ttttattgcg atctgcagat gaactggaaa atctcatttt acaacagaaac      60
tgagacagac gaccaccata ttcactgagg tctaaatttg cagtttccac taatgacatt      120
ttgatttccc aacagagata cttctggtct tactgcacag tcttttaaga gaaatacttc      180
cattatgcca cattgtcctt gatccgtaag tgatgtgtta aggtgcttca aaggaactct      240
gacctctgaa gtacttgagc tacttttagta tgtccagcct attgcttttt gtttttagtgt      300
gtcaccataa atatcagggg cataaaaaggc tatctattct taattcaagg ataaaaacaga      360
agaagcttgt ggtataaaac aatagttcaa gatccag                                     397

```

```

<210> 695
<211> 609
<212> DNA
<213> Homo sapien

```

```

<220>
<221> misc_feature
<222> (1)...(609)
<223> n = A,T,C or G

```

```

<400> 695
ctgagcttcc atttgtcagc tagcactgng gtagtcaacc atgcgaatga ggctattttg      60
gacctcatga ttgtccagtg cctgggctga taccngggga aacgaaattt tgtggctgcc      120
cacaaaatca tggaaaataa tgatttttta gaaaacctcc actgntttgt tgtgcagcaa      180
taaataactg aaacaccaat ccaaaaaact tataaagcta taacaattaa aacagnataa      240
taatagtncc gggatacaaa aatgggtcaaa ttgaagagga tacaaagcct caaagcagtc      300
ctcactcata ananccttgt tgtatcacta aaanggcatt aaaattgaga anaaggaana      360
actagtggat taattaataa atgagaagta tccataagga aaaattaataa ttnnattctt      420
gcttcacatt atgaaaaaat acaaacaaca gattgattaa agacttaaat gngatcaaca      480
aaatgttaaa actgtgataa gaacatttaa gaaaatagtt ctatnaccct gggataaaac      540
attttcntcc aaggcattaa agtgttaaat gaaaagactg atncatttat tcattagaat      600
ttaaattcn                                     609

```

```

<210> 696
<211> 300
<212> DNA
<213> Homo sapien

```

```

<400> 696

```

```
<210> 697
<211> 391
<212> DNA
<213> Homo sapien
```

<400> 697

```
<210> 698
<211> 536
<212> DNA
<213> Homo sapien
```

<400> 698

```
<210> 699
<211> 419
<212> DNA
<213> Homo sapien
```

```
<220>
<221> misc_feature
<222> (1)...(419)
```

<223> n = A,T,C or G

<400> 699

```
ngtccacctg agggcaggtg acaaggacct gacagagccc atgcagggct ttagatttgg      60
acacacaaga gttgataact tcctcatgaa ctccctgcct gatctaaact catattatgg     120
gttctgactg tttgagtaat catcttcaag gttaaaccctc ttggcagtta cccttttcac     180
aaagtgcaca gtgggaatcg agaatcgata ggggttaattt tggagcagtg gcttatacca     240
ttcacctctg tttttttgtg attatttcac agataatgag accttaataa caaataggcg     300
taaaaaaatt ttcacattga aatgatagaa acatttgatg taataaaaact tggttggctt     360
gatattttta ggaattgaaa cctagcaatc ttattggaga gacaagaatt ggtctccag     419
```

<210> 700

<211> 336

<212> DNA

<213> Homo sapien

<400> 700

```
ccacttattg tccttaaaaa tccatactga tacatggaca gtaagtgtgt tttcagatgg      60
agtaccagca ccgaaaatgg gttgagggag gatgggttgt atgtatgttt ctgccacta     120
attttgagca gccatattat gaattaaatc gtcacagcca agtaataacc caagaatggg     180
atgagtttca tgtgtaatag ctcaaagga ataagcatga atgctggagt ggaccattat     240
cctcaaatat tctatgtcac ttctcattta aagactcttg ttatgaacta ttagaaactt     300
taggcaaaat caaaagtatt tgcggcaaaa taaagg                                336
```

<210> 701

<211> 418

<212> DNA

<213> Homo sapien

<400> 701

```
ccatgtgatg atgttgacaa cccctgaaga gcctcagtc attgttccac gtttaagaac      60
taggaatacc aggactgatg caattctact gggtcactat cgcttggtcac aagacacaga     120
caatcagacc aaagtatttg ctgtaataac taagaaaaaa gaagaaaaac cacttgacta     180
taaatacaga tattttcgtc gtgtccctgt acaagaagca gatcagagtt ttcattgtggg     240
gctacagcta tgttccagtg gtcaccagag gttcaacaaa ctcatctgga tacatcatte     300
ttgtcacatt acttacaat caactgggtga gactgcagtc agtgcttttg agattgacaa     360
gatgtacacc cccttgttct tcgccagagt aaggagctac acagctttct cagaaagg     418
```

<210> 702

<211> 261

<212> DNA

<213> Homo sapien

<220>

<221> misc_feature

<222> (1)...(261)

<223> n = A,T,C or G

<400> 702

```
gggcctgttg tgggggtggg ggaagcaggg aggggaacag ctaaaatagg tgcgtgtgat      60
ttgggttaaaa aatagtaggg ggatgatgct aataattagg ctgnnggtgg ttgtgttgat     120
tcaaattatg tgttttttgg agagtcagtg cagtggtaga aatataattg ttgggacnat     180
tagnttttagc attggagtag gtttaggtta tgtacgtagt ctaggccata tgtgttggan     240
attgagacta gtagggctag g                                261
```

<210> 703
 <211> 261
 <212> DNA
 <213> Homo sapien

<220>
 <221> misc_feature
 <222> (1)...(261)
 <223> n = A,T,C or G

<400> 703
 gggcctgttg tgggggtggg ggaagcaggg aggggaacan ctaaataggt tgctgttgat 60
 ttggttaaaa aatagtaggg ggatgatgct aataattagg ctgnggggtgg ttgtgttgat 120
 tcaaattatg tgttttttgg agagtcatgt cagtggtagt aatataattg ttgggacnat 180
 tagnttttagc attggagtag gtttaggtta tgtacgtagn ctaggccata tgtgttggag 240
 attganacta gtagggctag g 261

<210> 704
 <211> 381
 <212> DNA
 <213> Homo sapien

<220>
 <221> misc_feature
 <222> (1)...(381)
 <223> n = A,T,C or G

<400> 704
 ngntntgaatt ctattaaaga tacaagagg agctggtacc atttcttctg aaactattac 60
 aaacaactga aaaggtggaa tttctcccta attcatttta ggaggccagc attatactga 120
 taccaaaacc tggcagaggt acaataataa aaggaaactt caagtcagta tcaactgatga 180
 acaccaatgt gaaaatcctc aataaaatac tggcaaaactg aattcagcag cacatcaaaa 240
 agctaatacca ccacaatcaa gtcagcttca tccctgcgat gcaagtctgg ttcaacatat 300
 gcaaataaat aaatacaatt catcagataa acagagctaa agacaaaatt cacatgattt 360
 tctcaataga tgcagaaaag g 381

<210> 705
 <211> 477
 <212> DNA
 <213> Homo sapien

<400> 705
 ctgaaccctc gtggagccat tcatacaggt ccctaattaa ggaacaagtg attatgctac 60
 ctttgcacgg ttagggtagc gcggccgtta aacatgtgtc actgggcagg cgggtgcctct 120
 aatactggtg atgctagagg tgatgttttt ggtaaacagg cggggtaaga tttgccgagt 180
 tccttttact ttttttaacc tttccttatg agcatgcctg tgttgggttg acagtgaggg 240
 taataatgac ttgttggtga ttgtagatat tgggctgtta attgtcagtt cagtgtttta 300
 atctgacgca ggcttatgcy gaggagaatg ttttcattgt acttatacta acattagttc 360
 ttctataggg tgatagattg gtccaattgg gtgtgaggag ttcagttata tgtttgggat 420
 ttttttaggtg gtgggtgttg agcttgaacg ctttcttaat tgggtggctgc ttttagg 477

<210> 706
 <211> 266

```
<220>  
<221> misc_feature  
<222> (1)...(266)  
<223> n = A,T,C or G
```

```
<210> 707
<211> 358
<212> DNA
<213> Homo sapien
```

```
<220>
<221> misc_feature
<222> (1)...(358)
<223> n = A,T,C or G
```

```
<210> 708
<211> 491
<212> DNA
<213> Homo sapien
```

```
<220>
<221> misc_feature
<222> (1)...(491)
<223> n = A,T,C or G
```

<400> 708						
cctactatgg	gngttaaatt	ttttactctc	tctacaaggt	tttttcctag	tgtccaaaga	60
gctgttcctc	tttggactaa	cagttaaatt	tacaagggga	tttagagggt	tctgtgggca	120
aattttaaagt	tgaactaaga	ttctatcttg	gacaaccagc	tatcaccagg	ctcggtaggt	180
ttgtgcctc	tacctataaa	tcttccact	atthtgcctac	atagacgggt	gtgctctttt	240
agctgttctt	aggtagctcg	tctggtttctg	ggggctcttag	ctttggctct	ccttgcaaag	300
ttatttctag	ttaattcatt	atgcagaagg	tataggggtt	agtccttgct	atattatgct	360
tggttataat	ttttcatctt	tcccttgcg	tactatatct	attgcgccag	gtttcaattt	420
ctatcgcta	tactttattt	gggtaaatgg	tttggctaag	gttgctctgg	agtaagggng	480
gaqtgggttt	g					491

<210> 709
 <211> 460
 <212> DNA
 <213> Homo sapien

<220>
 <221> misc_feature
 <222> (1)...(460)
 <223> n = A,T,C or G

<400> 709
 nggttttttt ttagagcaa ataatttatg caaaatatgt tacaaaatct gggatgctaa 60
 atagttgaca caagtactgt gtttgacatt tagtttcatt tgaattagta atagaatttg 120
 ctcttccaa catttacatc ttttttcttt ctgactttat atattttcaa taaaaatttg 180
 ctccacagtt ttttaagntca ttcttcttga atccgntttt acatttgctg ngacaaacct 240
 gcataaaact agattttata gatataactt ctttggaaga gataaaaatt caaaagtttg 300
 acattgcttt canttattct tttcttcatt gttttgattg gccctgttta gattgatgta 360
 ttgccaatct acttttgatg gcatgaatnt aaaatgacaa cataaaaagc ncttctagtg 420
 caacagtaat tgaaacttgc agttttccat taaaaaaaaa 460

<210> 710
 <211> 542
 <212> DNA
 <213> Homo sapien

<220>
 <221> misc_feature
 <222> (1)...(542)
 <223> n = A,T,C or G

<400> 710
 ctgttacagt gacaagagat aaaaagatag acctgcagaa aaaacaaact caaagaaatg 60
 tgttcagatg taatgtaatt ggagtgaata actgtgggaa aagtggagtt cttcaggctc 120
 ttcttggaag aaacttaatg aggcagaaga aaattcgtga agatcataga tctactatg 180
 cgattaacac tgtttatgta tatggacaag agaaatactt gttgttgcat gatattctag 240
 aatcggaatt tctaactgaa gctgaaatca tttgngatgt tgtatgcctg gtatataatg 300
 tcagcaatcc caaatccttt gaatactgtg ccaggatttt taagcaacac tttatggaca 360
 gcagaatacc ttgcttaatc gtagctgcaa agtcagacct gcatgaagtt aaacaagaat 420
 acagtatttc acctactgat ttctgcagga aacacaaaat gcctccacca caagccttca 480
 cttgcaatac tgctgatgcc cccagtnagg atatctttgt taaattgaca acaatggacc 540
 tg 542

<210> 711
 <211> 394
 <212> DNA
 <213> Homo sapien

<220>
 <221> misc_feature
 <222> (1)...(394)
 <223> n = A,T,C or G

<400> 711
 caaacccact ccaccttact accagacaac cttagccaaa ccatttacct aaataaagta 60

```

taggcgatag aaattgaaac ctggcgcaat agatatagta ccgcaaggga aagatgaaaa 120
attataacca agcataatat agcaaggact aacccctata ctttctgcat aatgaattaa 180
ctanaaataa ctttgcaagg agagccaaag ctaagacccc cgaaaccaga cgagctacct 240
aagaacagct aaaagagcac acccgtctat gtagcaaaat agtgggaaga tttataggna 300
gaggcgacaa acctaccgag cctggtgata gctggttgtc caagatagaa tcttagttca 360
actttaaaatt tgcccacaga accctctaaa tccc 394

```

```

<210> 712
<211> 552
<212> DNA
<213> Homo sapien

```

```

<220>
<221> misc_feature
<222> (1)...(552)
<223> n = A,T,C or G

```

```

<400> 712
gaggctctgta naatgccagg ctcaaatttg tctttataat ttaataaccag aaatctttcc 60
cttgtgatgt ttctttcttt ctggattgcc tctatagcag gggatagcgg gggaggataa 120
ggcacatctt tgntgtactg agaaatttga ccacgcagga tgatgtggct gttctcattc 180
atctgcacag agaaaaataa tgataaaata tccctttcct atgtttactg attttatggc 240
tgccataatg gaagcctcct tgactattta atcctttctg tcaactaggt tcgatttttt 300
ttttaattta cctgtagtag gtatttaana attttaacta gctanaaata attacattcc 360
aaaggaacac caaggcaaat aaatggttg taatcagcaa aagaattaca ttagttgttg 420
ntgctactta ttagggggag aactgttttt ttttaaattt aaacaattta ataatctcaa 480
ctgcaaataa ttttagatgc agcaaaggac tatgtagncg ttaataacct atgttgatat 540
tttcataata tt 552

```

```

<210> 713
<211> 518
<212> DNA
<213> Homo sapien

```

```

<220>
<221> misc_feature
<222> (1)...(518)
<223> n = A,T,C or G

```

```

<400> 713
ccaaaaactg gaagcagctc actaaacaaa cagtggcata cccatagaac tgcatacttc 60
tcagcagtat gaaagaatga gctacttata taagcatcat tgataaacct caaaaaaaaa 120
atgccacatg aanaaaccca aagggganaa acataaaaaac tttatatgtc agtcatataa 180
aattctanaa aatgcaaact aatccatcnt aaaggaaaagt aaatcaacag ttgtctggag 240
gaccananag agcaggagga ganagattat taaagggtt aaagtaaatt tgggagtgcc 300
cttcnntttt taaatnctat gaaaatgaaa gttaaaggcnc atgcatgttg taaactaata 360
gtaacaaaca naatgggttg gagtgggttg ttgtctgggg acatcattac aaaatgtaag 420
ccagtttatn taaattttga aaagaccgtg gactctgatc tgactgatna atgttggaag 480
agataagtgt gctgcaaagt ggggaattaa taaaacag 518

```

```

<210> 714
<211> 281
<212> DNA
<213> Homo sapien

```

```
<210> 715
<211> 443
<212> DNA
<213> Homo sapien
```

```
<210> 716
<211> 639
<212> DNA
<213> Homo sapien
```

<400>	716						
naaaa	tgaagtacag	agtctgcata	gtaagcttac	agataccttg	gtatcaaaac		60
ttgga	gcaaagacta	atgcagttaa	tggaatcaga	gcagaaaagg	gtgaacaaag		120
tctct	acaaatgcag	gttcaggata	ttttggagca	gaatgagggt	ttgaaagctc		180
cagca	gttcattcc	cagatagcag	cccagacctc	cgcttcagtt	ctagcagaag		240
cataa	agtgattgca	gaaaaggata	agcagataaa	acagactgaa	gattcttttag		300
gaacg	tgatcgttta	acaagtaaa	aagaggaaat	taaggatata	cagaatatga		360
ttatt	aaaagctgaa	gtgcagaaat	tacaggccct	ggcaaatgag	caggctgtcg		420
catga	attggagaag	atgcaacaaa	gtgtttatgt	taaagatgat	aaaataagat		480
gaaga	gcaactacaa	catgaatttt	caaacnaaat	ggaagaattt	angattctaa		540
caaaa	caanagcatta	aaatcagaag	ttcagaagct	gcagactctt	gtttctgcac		600
taata	aggatgntgn	ggaacaaatg	gaaaaattg				639

```
<220>  
<221> misc_feature  
<222> (1)...(473)
```



```

agataagtat caggtctgac cccagtggaa aacaaagcca aacaaaactg aaccacaaaa 240
aaaaaggctg gtgttcacca aaaccaaact tgttcattta gataatttga aaaagctcca 300
tagaaaaggc gtgcagtact aagggaacaa tccatgtgat taatgnttnc attatgttca 360
tgtaanaagc cccttatttt tagccataat tttgcatact gaaaatccaa taatcagaaa 420
agtaattttg ccacattatt tatnaaaaat gttcc 455

```

```

<210> 721
<211> 530
<212> DNA
<213> Homo sapien

```

```

<220>
<221> misc_feature
<222> (1)...(530)
<223> n = A,T,C or G

```

```

<400> 721
ccagtgtttg ctgccgtggt ttagtgattg ggtgttagaa ataaaaactc aggtctatatt 60
cttaccagtc agtaacaatt tttagagaat gtacttggtg tataatatat ggacttcagg 120
aacttttatt gggngggggg ttaattttgc cttaccctgt tcactttcag atgattaggc 180
ttttgcactt tagaatgaga aacttgtgac gttagtgtgt tcttactagc ttttaatttgt 240
atgtagcaat gaattgtgaa tcttagtgca gtgggttttt ttaaaaaact caaaaagctg 300
ggaattaagt ggttttcagta ataatgctat accgaggtgc ttgcattgta tttcataatt 360
ttgttacaaa ccaaaattat ttttaatgan aacggtcctg ggttcagagg tgtgatgcc 420
gaatgtatatt tctgactgtt aggcccttgg aacagatacc ggtgctttct tgaaagatga 480
aagaaatgca atgggtgctc ttcatgcaag gttgcaaacc taccaagaat 530

```

```

<210> 722
<211> 242
<212> DNA
<213> Homo sapien

```

```

<220>
<221> misc_feature
<222> (1)...(242)
<223> n = A,T,C or G

```

```

<400> 722
ccaaggggtca tgatggcagg agtaatcana ggtgntcctg tgttgtgata agggngggaga 60
ggttaaagga gccacttatt agtaatgttg atagtagaat gatggctagg gtgacttcat 120
atgagattgt ttgggctact gctcgcagtg cgccgatcag ggcgtagttt gagtttgatg 180
ctcatcctga tnagaggatt gagtaaacgg ctaggctaga ggtggctaga ataaatagga 240
gg 242

```

```

<210> 723
<211> 472
<212> DNA
<213> Homo sapien

```

```

<220>
<221> misc_feature
<222> (1)...(472)
<223> n = A,T,C or G

```

```
<210> 724
<211> 292
<212> DNA
<213> Homo sapien

<220>
<221> misc_feature
<222> (1)...(292)
<223> n = A,T,C or G
```

```
<210> 725
<211> 122
<212> DNA
<213> Homo sapien

<220>
<221> misc_feature
<222> (1)...(122)
<223> n = A,T,C or G
```

```
<210> 726
<211> 477
<212> DNA
<213> Homo sapien

<220>
<221> misc_feature
<222> (1)...(477)
<223> n = A,T,C or G
```

<400> 726
ctgaacccctc gtggagccat tcatacaggt ccctaattaa ggaacaagtq attatgctac 60


```

<210> 730
<211> 310
<212> DNA
<213> Homo sapien

<400> 730
ccattttttat ttctttcttca gagaagtggt tatttaggtc tgttgcccat tttaacaatta      60
ggccatatgt tttcttgctg ttgagttgta tgtgtgtttg tataaatttt gcatattaac      120
cccttatcac acgtatgttt tttaaaaataa attttgctta ttaatctttt atcagatgta      180
tggttttcaa atatattctt ccgatccatg gattctcttt tttgttatga ttgtttcttt      240
gctcttcgga agctttttgt tttgttttgt tatttgtttt actttgatat agtcccattt      300
attgtttttg                                     310

<210> 731
<211> 467
<212> DNA
<213> Homo sapien

<220>
<221> misc_feature
<222> (1)...(467)
<223> n = A,T,C or G

<400> 731
ngacaacctt agccaaacca tttacccaaa taaagtatag gcgatagaaa ttgaaacctg      60
gcgcaataga tatagtaccg caagggaaaag atgaaaaatt ataaccaagc ataataaagc      120
aaggactaac ccctatacct tctgcataat gaattaacta gaaataactt tgcaaggaga      180
gccaaagcta agacccccga aaccagacga gctacctaag aacagctaaa agagcacacc      240
cgtctatgta gcaaaatagn gggaagattt atagnnagag gcgacaaacc taccgagcct      300
ggtgatagct gggtgtccaa gatagaatct tagntcaact ttaaatttgc ccacagaacc      360
ctctaaatcc ccttgtaaat ttaactgnta gnccaaagag gaacagntct ttggacacta      420
ggaaaaaacc ttgtagagag agtaaaaaat ttaacaccca tagtagg      467

<210> 732
<211> 492
<212> DNA
<213> Homo sapien

<220>
<221> misc_feature
<222> (1)...(492)
<223> n = A,T,C or G

<400> 732
cctactatgg gtgttaaatt ttttactctc tctacaaggt tttttcctag tgtccaaaga      60

```


atttaatacg tgctcactgc tcggcacgcg ctgaagctac agttaacaat cagtgagcac 60
 atattaaatg ataaaataat gctgatggta aacattcata acagcagagt aagattttgg 120
 cagttttgtg tctcggtaac ataactgtaa ccttagatga acacctatcc cttcatgac 180
 tgactttaga ggcaaggagt ttgtaacatc taatgg 216

<210> 736

<211> 285

<212> DNA

<213> Homo sapien

<220>

<221> misc_feature

<222> (1)...(285)

<223> n = A,T,C or G

<400> 736

ctgaaaggca acntggagac tagttagtct agtccccctca tattataaat tggatatgctg 60
 aggccaggca gtaaatgtct atggagctct ccaatttaag gccagtttga ctccaagggt 120
 agggcttcta gtaaaatttt gtgattaaat tggaaaactct aatttatattt tctatgngtt 180
 tttggtacct aatcctcata agcaagccat atttcaaggc tgatcaatga aaacacccaaa 240
 taccaaagct tcctttccct tccaaattta ctgacccttt gtcag 285

<210> 737

<211> 509

<212> DNA

<213> Homo sapien

<220>

<221> misc_feature

<222> (1)...(509)

<223> n = A,T,C or G

<400> 737

agangaagaa gangaagatt aagggaaaag tacatcggtc aagaagagct caacaaaaca 60
 aagcccctct ggaccagaaa tcccgcgat attactaatg aggagtaagg agaattctat 120
 aagagcttga ccaatgactg ggaagatcac ttggcagtga agcatttttc agttgaagga 180
 cagttggaat tcagagccct tctatttgtc ccacgcagtg ctccctttga tctgtttgaa 240
 aacagaaaga aaaagaacaa catcaaattg tatgtacgca gagttttcat catggataac 300
 tngaggagc taatccctga atatctgaac ttcattagag ggggtggnaga ctggaggat 360
 ctccctctaa acatatcccg tgagatgttg caacaaagca aaattttgaa agttatcang 420
 aagaatttgg gtcaaaaaat gcttanaact ctttactgaa ctggcggaag atnaagagaa 480
 ctncagana ttctatgagc agntctctt 509

<210> 738

<211> 97

<212> DNA

<213> Homo sapien

<400> 738

cagtgaattg aatacgactc ctatagggcg aattgggccc tctagatgca tgctcgagcg 60
 gccgccagtg tgatggatat ctgcagaatt cgccctt 97

<210> 739

<211> 209

<400> 745
 cgcgttactg tacatattgc tagcaggaga caactggaaa tactaaacaa atactggaat 60
 tcacattaca gacagacgaa accaacatgg atgccacaca taacttcctt tgtagtttca 120
 cagagagcct atttgtggtt gctcagggtg ggtcatacat tgcttgcaga aatggcctga 180
 tcatagctct atgaaacaat gaattcggaa tgaaatctta ccatgacacc tctctgtagg 240
 aaagaaatgt tgcttcacgt gtgctaagtt gagataataa tatttcacat atttatatac 300
 agagaatcac tctcaaattt aacccaagat aagcaatagg atttgggggt gacttgtaga 360
 catttctaac aacacttttc ttttttctag aggtcactct caaactga tatatcata 420
 tagtttgagt gtanggattc agtaatcaaa gggtgttatt gcaaaagagc caggcag 477

<210> 746
 <211> 524
 <212> DNA
 <213> Homo sapien

<220>
 <221> misc_feature
 <222> (1)...(524)
 <223> n = A,T,C or G

<400> 746
 ctgtgaaatt ggggtgggag agccaaaata ctttacaact tcagaccgga gaaaaggcca 60
 gaggtgtgaa gttagactct atgatgaaac agagtctgtt tttgcatga catgttggga 120
 taatgaatcc attctacttg cacagagctg gatgccacga gaaacagtaa tatttgcctc 180
 agatgtaaga ataaattttg acaaatttcg gaactgcatg acagcaactg taatctcaaa 240
 aaccattatt acaactaatc cagatatacc agaagctaac attctgctga attttatacg 300
 agaaaataaa gaaacaaatg ttctggatga tgaaattgac agttatttca aagaatccat 360
 aaatttaagt acaatagttg atgtctacac agntgaacaa ttaaaggga aagctttgaa 420
 gaatgaagga aaagctgatc cttcctatgg catcctttat gctacattt ccacactcaa 480
 cattgatgat gaaactcaaa agtagttcga aatagatggt ccag 524

<210> 747
 <211> 456
 <212> DNA
 <213> Homo sapien

<220>
 <221> misc_feature
 <222> (1)...(456)
 <223> n = A,T,C or G

<400> 747
 cctcagttct tgattgtggt tgacggggcg tcaccatgaa ggagcccatt tagtataaag 60
 cttccaacct tttctcttaa tcgtttcttt aatcttttaa accatcttca agtgcatagg 120
 ggagtttccg atgccagagg atgaaagcaa gtgctttctc caccctctcc tcccagagtg 180
 aaaacaaatc cttttgctga tacttgtttc aaaagcatcc attgtaaagc ttctcagtga 240
 cacaaaatac tgagaggtaa ctttttatca atcaaacac ataccccaat ttaacacctt 300
 tcagtgtctc gaattcaact gacagactaa aggggtgttc ctgtaacagt ctgaaatatt 360
 aagtgttttt tttgttttgt ttttaaatct ttttcagaa aacttcctct nggggtagga 420
 aagtacacat gaagcagcaa agtaacgaag aaaaac 456

<210> 748
 <211> 474

<212> DNA
<213> Homo sapien

<220>
<221> misc_feature
<222> (1)...(474)
<223> n = A,T,C or G

```

<400> 748
ccanaccagg gaaccaaaatg cagacagnga agttctctgc ttcttttggc tataatgnga      60
caagaaaggg atcatctttt gaagatgttt aaagaaataa agcaactttc ttataaaaca      120
gtcaaataat caattaatgg aataaataag tactaaccca cattttaacc actctgtaat      180
cactacactt tacatathtt ttatttnggn ggcaaantcc ccataatta gtctaaaatc      240
caccaatcac ttttaaaagt aaaatgaata gccaccaaaa taagaaaatc ttctgttcac      300
tctttggcta aaaaggaaaa caaataaaac aaaacaaaaa gaaacagaag acaactgtaa      360
cactgggtgat aaaagaaact ttttttttac aagtaaaata aagttatcaa tttaaatctt      420
ggncacttta taaaaacaag aggtaatgtt gtaataaaac agcagtagcc tcag              474

```

<210> 749
<211> 355
<212> DNA
<213> Homo sapien

<220>
<221> misc_feature
<222> (1)...(355)
<223> n = A,T,C or G

```

<400> 749
cctgggttna gnggctgact gnaacctcca cttcctgttc tcaggcaatc ctctgcctc      60
agcctcctta gtagctggga ctacaggagt gtgcaaccat gcccaactaa tttttgtatt      120
tttaatatag acagggtttc accatgttga tcaggttggg ctccaactcc tgacctcagg      180
tgatccacct gtcccagcct cccaaagtgc tgggattaca ggcatgagcc accacgcccg      240
gnccaggata aagtaaaaat ttgtaagcac acaaggccct ttgcaacctg gctcctgggt      300
actactttta ncctcctgcc ctcccaaatg tinctactgt ttttctanac atacc              355

```

<210> 750
<211> 493
<212> DNA
<213> Homo sapien

<220>
<221> misc_feature
<222> (1)...(493)
<223> n = A,T,C or G

```

<400> 750
ccatgtctgg ctcgaactcc tgaactcagg tgatccaccc gcctcagtct cccaatagat      60
tacatatatt attaatgaat tgcttccttt aacaccctat tcattgaatt ttccagtaaa      120
ccacaattac taattactcc tgaatcaga aaagagggtta aaaagatttt ataacagtat      180
cctatgaaat ctactacttt caagtaatag tagttgaatt accaaaaacc gtcactcaag      240
ccaatgacta caattaagat atgagtaaca tttcctagat aaataaagtc aattaattat      300
atgtgcatct gggaaataga gaaagtacat ataagccatg attttgaagn caaaagagag      360
agantatttg ccaaggaggg gtgagttata gtatgtaatt ataacataca gaagcttttt      420

```



```

nacaacctta gccanaacca tttacccaaa taaagggata ggcgatagaa attgaaacct      60
ggcgcaatag atatagnacc gcaagggaaa gatgaaaaat tataaccaag cataatatag      120
caaggactaa cccctatacc ttctgcataa tgaattaact agaaataact ttgcaaggag      180
agccaaaagct aagacccccg aaaccagacg agctatctaa gaacagctaa aagagcacac      240
ccgtctatgt agcaaaatag tgggaagatt tataggtaga ggcgacaaac ctaccgagcc      300
tgggtgatagc tggntgncca agatagaatc ttagntcaac tttaaatttg cccacagaac      360
cctctaaatc cccttgtaaa ttttaactgtt agtccaaaga ggaacagctc ttggacacna      420
ggaaaaaacc ttgcagagag agtaaaaaat ttaacacca tagtagg                      467

```

```

<210> 754
<211> 196
<212> DNA
<213> Homo sapien

```

```

<220>
<221> misc_feature
<222> (1)...(196)
<223> n = A,T,C or G

```

```

<400> 754
gtcatgttca agtgttntaa tctgacgcag gcttatgcgg aggagaatgt tttcatgtta      60
cttatactaa cattagtctt tctatagggt gatagattgg tccaattggg tgtgaggagt      120
tcagttatat gtttgggatt ttttaggcag tgggtggtga gcttgaacgc tttcttaatt      180
ggtggctgct tttagg                      196

```

```

<210> 755
<211> 381
<212> DNA
<213> Homo sapien

```

```

<400> 755
ctggaaagga ttctgtacat ataagacatc aaatattgag ggatactgga actttttaaat      60
taatgggcaa agaaagtcaa caaaggaagt tcatatgaaa tcaaactagt aatatgatta      120
caaaaaaaaaa gtttaaaatt tttcttggcc ccagtcttat catttctgag ccaaatacaa      180
ttctatcgaa atcacctgaa actgaaatca ccattctagg ctgggtttcc cataaagatg      240
gactgtctcca aaaagaggaa tcaagaaaaga atttggctca cagtgaatta ttcaactttgt      300
cttagttaac taaaaataaa atctgactgt taactacaga aatcatttca aattctgtgg      360
tgataataaa gtaatgaccg c                      381

```

```

<210> 756
<211> 341
<212> DNA
<213> Homo sapien

```

```

<220>
<221> misc_feature
<222> (1)...(341)
<223> n = A,T,C or G

```

```

<400> 756
ggntataaac ctattattta ttgcagaact aataaaaaat ccaaagcctt gtatttgtac      60
atctttatta tctctaaagc actttcctca acctaatttc agtttttaca attggtactc      120
aagaaaaatag agacagaaat catttgattt tgcccagaaa ccatctgctt atatttataa      180
ggccacctaa tttgaaatca catatagacc aggcgcggtg gctcacgcct gtaattccaa      240

```



```

ttgaaccagn ccagtctgat tttcaggtga attctgtgaa gagcttgatg ggggaagtct 240
gaagacagaa ggaattaggg aaaaggggtga tacttacaga gtaaaggaaa taaatgaaaa 300
gataatggta tttttggtag ccacagggaa atagcaggag gggactggag atcacacaca 360
cgcacacgca cacacacaaa cacacacaca cgctaaaact caaactaaaa acctcccaa 420
ggagctgctt tgtttgcaga cttcaattng aagtagatac taagggaag aatagaccag 480
ttaaattca cctgaaaatc tcttcccann cttcaaatgt gctaaaatat cactgtcagc 540
ttagcatctc tncatgtatg tatatataga tgta 574

```

```

<210> 763
<211> 465
<212> DNA
<213> Homo sapien

```

```

<220>
<221> misc_feature
<222> (1)...(465)
<223> n = A,T,C or G

```

```

<400> 763
cctactatgg gtgttaaaat tttttactct ctctacaagg ntttttcta gtgtccaaag 60
agctgttctt ctttggaacta acagttaaat ttacaagggg atttagaggg ttctgngggc 120
aaatttaaaag ttgaactaag attctatctt ggacaaccag ctatcaccag gctcggtagg 180
tttgctgcct ctacctataa atcttcccac tattttgcta catagacggg tgtgctcttt 240
tagctgttct taggtagctc gtctgggttc ggggggtctta gctttggctc tccttgcaaa 300
gttatttcta gttaattcat tatgcagaag gtataggggt tagtccttgc tatattatgc 360
ttggatataa tttttcatct ttcccttgcg gtactatatac tattgcgcca ngtttcaatt 420
tctatcgctt atactttatt tgggtaaatg gtttggtctaa ggttg 465

```

```

<210> 764
<211> 151
<212> DNA
<213> Homo sapien

```

```

<400> 764
ctgtcaatta atgctagtcc tcaggattta aaaaataatc ttaactcaaa gtccaatgca 60
aaaacattaa gttggtaatt actcttgatc ttgaattact tccggtacga aagtccttca 120
catttttcaa actaagctac tatatttaag g 151

```

```

<210> 765
<211> 251
<212> DNA
<213> Homo sapien

```

```

<400> 765
gaagagctta tcacctttca tgatcacgcc ctcatagtca ttttccttat ctgcttcta 60
gtcctgtatg cctttttcct aacactcaca acaaaaactaa ctaataactaa catctcagac 120
gctcaggaaa tagtaaccgt ctgaactatc ctgcccgcga tcatcctagt cctcatcgcc 180
ctcccatccc tacgcatcct ttacataaca gacgaggtca acgatccctc ccttaccatc 240
aatcaattg g 251

```

```

<210> 766
<211> 375
<212> DNA
<213> Homo sapien

```

<220>
 <221> misc_feature
 <222> (1)...(375)
 <223> n = A,T,C or G

<400> 766
 cgaggtctgn cctcctgggt cttcatccat tattaacaga agagcatact gggttcgggc 60
 cataaaatct ttgggaaggg acaactgtaa aggaagttca tagtcgtcaa tatgaaggat 120
 ttttaatttct ggctttccta tcttcttctt caggatagct tccttcagca tagaattgtt 180
 ttccaatata aaatattttg ctgggttggt cgtactatgt aggctgacca ctgggaccct 240
 tggaccttca cagaataata agaaatgttg attcatggga ctaaaactgg catcaaaata 300
 tgtacattgt tctttcatga aattacatga aatgcattgg cgattcaata atccttcagt 360
 agaagcactg tacag 375

<210> 767
 <211> 485
 <212> DNA
 <213> Homo sapien

<220>
 <221> misc_feature
 <222> (1)...(485)
 <223> n = A,T,C or G

<400> 767
 cgaggtctga accctcgtgg agccattcat acaggtccct aattaaggaa caagtgatta 60
 tgctaccttn gcacgggttag ggtaccgcgg cccgttaaac atgtgtcact gggcaggcgg 120
 tgcctctaact actggtgatg ctagaggtga tgtttttggn aaacaggcgg ggtaagattt 180
 gccgagttcc ttttactttt tttaaccttt ccttatgagc atgcctgtgt tgggttgaca 240
 gtgagggtaa taatgacttg ttggtgattg tagatattgg gctgttaatt gtcagttcag 300
 tgttttaate tgacgcaggc ttatgcggag gagaatgttt tcatgttact tatactaaca 360
 ttagttcttc tatagggtga tagatnggtc caattgggtg tgaggagntc acttatatgt 420
 ttgggatttt ttaggtaagn ggggtgtgag cttgaacgct ttcttaattg ggggctgctt 480
 ttang 485

<210> 768
 <211> 379
 <212> DNA
 <213> Homo sapien

<220>
 <221> misc_feature
 <222> (1)...(379)
 <223> n = A,T,C or G

<400> 768
 ctgatattct attaaagata caaagaggag ctggnacat ttcttctgaa actattacaa 60
 acaactgaaa aggtggaatt tctccctaact tcatttttagg aggccagcat tatactgata 120
 ccaaaacctg gcagaggtac aataataaaa ggaaacttca agtcagtatc actgatgaac 180
 accaatgtga aaatcctcaa taaaatactg gcaaactgaa ttcagcagca catcaaaaag 240
 ctaatccacc acaatcaagt cagcttcac cctgcgatgc aagtctgggt caacatatgc 300
 aaatcaataa atacaattca tcagataaac agagctaaag acaaaattca catgattttc 360
 tcaatagatg cagaaaagg 379

<210> 769
 <211> 518
 <212> DNA
 <213> Homo sapien

<220>
 <221> misc_feature
 <222> (1)...(518)
 <223> n = A,T,C or G

<400> 769
 cgagggtccat atgatgatca gtctatatag ttttaaggcgc agatacacaa attttcacaaa 60
 atatgggtag aatatagtca atatgaatgg aatagacaat gctttgaaaa tccactggagg 120
 gaggcctttat tgttttgtgaa aacatgttgt catcactttt tgctttaagc ccttggtggt 180
 gaaataactc aaaccattct tccttatgct gaagatcgag aaccccaagt atcacatcta 240
 ccatcccact catcaatgtg attggtcagt ctttgcagag gncctgcata gccagtttta 300
 aagtttagagt tcttgcata acatatgaaa aggcattgta cttgtgcttt caaagagctt 360
 tttgcttgggt gtaaaaagaa aactcaaatt acagtgtgat gtggaatata atgggtgtag 420
 tttcatcgag atgatgggaa agaattgata agataaagcn gaaagatgag cagaattttc 480
 agattgggtn tggaaagagc acttaagaaa gaggggtgg 518

<210> 770
 <211> 378
 <212> DNA
 <213> Homo sapien

<220>
 <221> misc_feature
 <222> (1)...(378)
 <223> n = A,T,C or G

<400> 770
 tatgggtcct gagtgtggaa tataagataa caagacaatt cccttgcttt caagggaaat 60
 cacactttat aaaactttga attcctgaaa tgggtttcag aggttccaag gtcaaattca 120
 agaataagag ttaagaagaa aaagactatg agaaaggaag tgntgacccc atttgcattt 180
 aaatggcagg aatagtctca atctactcat tggggaaaaa tgtatgttgc atatttttga 240
 gatattgcaa cttgctctct ctctttgcca ccccaccctt tgnatgctc tgtttttggg 300
 ctgaattggc aagaaaaatg gctggagggc tggagaaggn tggacccttc ttccttcttc 360
 cttcttcttc ctttctcc 378

<210> 771
 <211> 207
 <212> DNA
 <213> Homo sapien

<400> 771
 cataaatatt atactagcat ttaccatctc acttctagga atactagtat atcgctcaca 60
 cctcatatcc tccctactat gcctagaagg aataatacta tccactgttca ttatagctac 120
 tctcataacc ctcaacaccc actcctctt agccaatatt gtgcctattg ccatactagt 180
 ctttgcogcc tgcgaagcag cggtagg 207

<210> 772
 <211> 384

<213> Homo sapien

<220>

<221> misc_feature

<222> (1)...(277)

<223> n = A,T,C or G

<400> 779

cctnttgatt	tgatgggtaa	ggggagggat	cgttgacctc	gtctgttatg	taaaggatgc	60
gtagggatgg	gagggcgatg	aggactagga	tgatggcggg	caggatagtt	cagacggttt	120
ctatttcctg	agcgtctgag	atgttagtat	tagttagttt	tgttgtgagt	gttaggaaaa	180
gggcatacag	gactaggaag	cagataagga	aaatgactat	gagggcgtga	tcatgaaagg	240
tgataagctc	ttctatgata	ggggaagtag	cgtcttg			277

<210> 780

<211> 328

<212> DNA

<213> Homo sapien

<220>

<221> misc_feature

<222> (1)...(328)

<223> n = A,T,C or G

<400> 780

catgntatgg	ataacatnt	taactgtatt	ttntgcanc	cgtaccttct	tgggaataca	60
attgtctaac	tttttat	tggnctggct	gttgtgggtg	gcaaaactcc	gtacattgct	120
atthttgccac	actgcaacac	cttacagatg	tggaagatgt	gaaatttgtc	atcaattatg	180
actaccctaa	ctcctcagag	gattatattc	atcgaattgg	aagaactgct	cgcagtacca	240
aaacaggcac	agcatacact	ttcttttacac	ctaataacat	aaagcagggg	agcgacctta	300
tctctgtgct	tcgggaagct	aancaaac				328

<210> 781

<211> 305

<212> DNA

<213> Homo sapien

<220>

<221> misc_feature

<222> (1)...(305)

<223> n = A,T,C or G

<400> 781

ctgttcagaa	agctcattgg	acctggtttt	gaaaataaaa	caaagttaaa	accctgggag	60
gagttattgt	gcagngtgga	gtactcaggc	tttcttataa	agaaaaaaaa	agttatctgg	120
taccaaagtg	tgcaacctac	agaccctcag	gtactgccct	gtgacttctc	tgtatgacat	180
cacaaggctg	ccaagtgcct	gtttttctag	aactaggagt	tggtgagggt	tggctantgc	240
tgaaaccatg	cataggattg	gtttactaaa	ttaaaacctt	attacgtacg	tcctccaaaa	300
gacag						305

<210> 782

<211> 497

<212> DNA

<213> Homo sapien

<220>
 <221> misc_feature
 <222> (1)...(497)
 <223> n = A,T,C or G

<400> 782
 cgaggtggct ttaattgatg ttaatgcctt atgtcaaattg taaagttaga atttgctagg 60
 gctgggatag ggagtgatat ttctaggact tagacattga aaactaattc agcctgtagt 120
 aacctggatg gttttcaatg gcatggttag tcaaattcat ggtttttaaac ttagaagcag 180
 ctttcggggg agagggtagg ttggagcatt tattacatat ttactgttt aatgtcttaa 240
 ccgtgggcct ttttaatttgt aaacactgaa atgattgttg ggctgtggaa aacatttacc 300
 tatttacctt ggaagtttta aaagacagtc cacttttttag catgtgtgtt gcgtccagcc 360
 tgtggtcgtc ttaactaata aatgngattt ttctctcaa aaaaaaacct ccccgggcgg 420
 ccgtcaagg gnaattccn cacactggcg gccgttacta ggggatccga nctcgggtcca 480
 agcttggcgt aatcatg 497

<210> 783
 <211> 364
 <212> PRT
 <213> Homo sapien

<400> 783
 Met Trp Gln Pro Leu Phe Phe Lys Trp Leu Leu Ser Cys Cys Pro Gly
 1 5 10 15
 Ser Ser Gln Ile Ala Ala Ala Ala Ser Thr Gln Pro Glu Asp Asp Ile
 20 25 30
 Asn Thr Gln Arg Lys Lys Ser Gln Glu Lys Met Arg Glu Val Thr Asp
 35 40 45
 Ser Pro Gly Arg Pro Arg Glu Leu Thr Ile Pro Gln Thr Ser Ser His
 50 55 60
 Gly Ala Asn Arg Phe Val Pro Lys Ser Lys Ala Leu Glu Ala Val Lys
 65 70 75 80
 Leu Ala Ile Glu Ala Gly Phe His His Ile Asp Ser Ala His Val Tyr
 85 90 95
 Asn Asn Glu Glu Gln Val Gly Leu Ala Ile Arg Ser Lys Ile Ala Asp
 100 105 110
 Gly Ser Val Lys Arg Glu Asp Ile Phe Tyr Thr Ser Lys Leu Trp Ser
 115 120 125
 Asn Ser His Arg Pro Glu Leu Val Arg Pro Ala Leu Glu Arg Ser Leu
 130 135 140
 Lys Asn Leu Gln Leu Asp Tyr Val Asp Leu Tyr Leu Ile His Phe Pro
 145 150 155 160
 Val Ser Val Lys Pro Gly Glu Glu Val Ile Pro Lys Asp Glu Asn Gly
 165 170 175
 Lys Ile Leu Phe Asp Thr Val Asp Leu Cys Ala Thr Trp Glu Ala Met
 180 185 190
 Glu Lys Cys Lys Asp Ala Gly Leu Ala Lys Ser Ile Gly Val Ser Asn
 195 200 205
 Phe Asn His Arg Leu Leu Glu Met Ile Leu Asn Lys Pro Gly Leu Lys
 210 215 220
 Tyr Lys Pro Val Cys Asn Gln Val Glu Cys His Pro Tyr Phe Asn Gln
 225 230 235 240
 Arg Lys Leu Leu Asp Phe Cys Lys Ser Lys Asp Ile Val Leu Val Ala

245 250 255
 Tyr Ser Ala Leu Gly Ser His Arg Glu Glu Pro Trp Val Asp Pro Asn
 260 265 270
 Ser Pro Val Leu Leu Glu Asp Pro Val Leu Cys Ala Leu Ala Lys Lys
 275 280 285
 His Lys Arg Thr Pro Ala Leu Ile Ala Leu Arg Tyr Gln Leu Gln Arg
 290 295 300
 Gly Val Val Val Leu Ala Lys Ser Tyr Asn Glu Gln Arg Ile Arg Gln
 305 310 315 320
 Asn Val Gln Val Phe Glu Phe Gln Leu Thr Ser Glu Glu Met Lys Ala
 325 330 335
 Ile Asp Gly Leu Asn Arg Asn Val Arg Tyr Leu Thr Leu Asp Ile Phe
 340 345 350
 Ala Gly Pro Pro Asn Tyr Pro Phe Ser Asp Glu Tyr
 355 360

<210> 784
 <211> 6353
 <212> DNA
 <213> Homo sapien

<400> 784
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 ctttctcgcc acgttcgccc gctttccccg tcaagctcta aatcgggggc tccctttagg 180
 gttccgattt agtgctttac ggcacctcga ccccaaaaaa cttgattagg gtgatggttc 240
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 ctttaaatag ggactcttgt tccaaactgg aacaacactc aaccctatct cgggtctattc 360
 ttttgattta taagggattt tgccgatttc ggctatttgg ttaaaaaatg agctgattta 420
 acaaaaaattt aacgcgaatt ttaacaaaat attaacgttt acaatttcag gtggcacttt 480
 tcggggaaat gtgcgcggaa cccctatttg tttatttttc taaatacatt caaatatgta 540
 tccgctcatg aattaattct tagaaaaact catcgagcat caaatgaaac tgcaatttat 600
 tcatatcagg attatcaata ccatattttt gaaaaagccg tttctgtaat gaaggagaaa 660
 actcaccgag gcagttccat aggatggcaa gatcctggta tcggtctgcg attccgactc 720
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<210> 785

<211> 5502

<212> DNA

<213> Homo sapien

<400> 785

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<210> 786

<211> 108

<212> PRT

<213> Homo sapiens

<400> 786

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Ala Ser Pro Arg Ser Pro Val Met Glu Ser Pro Lys Lys Lys Asn Gln
              35              40              45
Gln Leu Lys Val Gly Ile Leu His Leu Gly Ser Arg Gln Lys Lys Ile
              50              55              60
Arg Ile Gln Leu Arg Ser Gln Val Leu Gly Arg Glu Met Arg Asp Met
              65              70              75              80
Glu Gly Asp Leu Gln Glu Leu His Gln Ser Asn Thr Gly Asp Lys Ser
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Gly Phe Gly Phe Arg Arg Gln Gly Glu Asp Asn Thr
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<210> 787

<211> 152

<212> PRT

<213> Homo sapiens

<400> 787

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              20              25              30
Asn Asn Glu Glu Gln Val Gly Leu Ala Ile Arg Ser Lys Ile Ala Asp
              35              40              45
Gly Ser Val Lys Arg Glu Asp Ile Phe Tyr Thr Ser Lys Leu Trp Ser
              50              55              60
Thr Phe His Arg Pro Glu Leu Val Arg Pro Ala Leu Glu Asn Ser Leu
              65              70              75              80
Lys Lys Ala Gln Leu Asp Tyr Val Asp Leu Tyr Leu Ile His Ser Pro

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<210> 788
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<212> DNA
<213> Homo sapiens
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<210> 789
<211> 200
<212> PRT
<213> Homo sapien
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<400> 789
Met Ala Lys Gly Asp Pro Lys Lys Pro Lys Gly Lys Met Ser Ala Tyr
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 35 40 45
 Trp Lys Thr Met Ser Gly Lys Glu Lys Ser Lys Phe Asp Glu Met Ala
 50 55 60
 Lys Ala Asp Lys Val Arg Tyr Asp Arg Glu Met Lys Asp Tyr Gly Pro
 65 70 75 80
 Ala Lys Gly Gly Lys Lys Lys Lys Asp Pro Asn Ala Pro Lys Arg Pro
 85 90 95
 Pro Ser Gly Phe Phe Leu Phe Cys Ser Glu Phe Arg Pro Lys Ile Lys
 100 105 110
 Ser Thr Asn Pro Gly Ile Ser Ile Gly Asp Val Ala Lys Lys Leu Gly
 115 120 125
 Glu Met Trp Asn Asn Leu Asn Asp Ser Glu Lys Gln Pro Tyr Ile Thr
 130 135 140
 Lys Ala Ala Lys Leu Lys Glu Lys Tyr Glu Lys Asp Val Ala Asp Tyr
 145 150 155 160
 Lys Ser Lys Gly Lys Phe Asp Gly Ala Lys Gly Pro Ala Lys Val Ala
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<210> 790

<211> 457

<212> DNA

<213> Homo sapiens

<400> 790

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<210> 791

<211> 126

<212> PRT

<213> Homo sapiens

<400> 791

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Pro Glu Val Trp Ile Leu Ser Pro Leu Leu Arg His Gly Gly His Thr
 20 25 30

Gln Thr Gln Asn His Thr Ala Ser Pro Arg Ser Pro Val Met Glu Ser
 35 40 45

Pro Lys Lys Lys Asn Gln Gln Leu Lys Val Gly Ile Leu His Leu Gly
50 55 60

Ser Arg Gln Lys Lys Ile Arg Ile Gln Leu Arg Ser Gln Cys Ala Thr
65 70 75 80

Trp Lys Val Ile Cys Lys Ser Cys Ile Ser Gln Thr Pro Gly Ile Asn
85 90 95

Leu Asp Leu Gly Ser Gly Val Lys Val Lys Ile Ile Pro Lys Glu Glu
100 105 110

His Cys Lys Met Pro Glu Ala Gly Glu Glu Gln Pro Gln Val
115 120 125

<210> 792

<211> 461

<212> DNA

<213> Homo sapiens

<400> 792

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<210> 793

<211> 108

<212> PRT

<213> Homo sapiens

<400> 793

Arg Arg Ser Cys Glu Pro Ala Thr Arg Val Pro Glu Val Trp Ile Leu
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Ser Pro Leu Leu Arg His Gly Gly His Thr Gln Thr Gln Asn His Thr
20 25 30

Ala Ser Pro Arg Ser Pro Val Met Glu Ser Pro Lys Lys Lys Asn Gln
35 40 45

Gln Leu Lys Val Gly Ile Leu His Leu Gly Ser Arg Gln Lys Lys Ile
50 55 60

Arg Ile Gln Leu Arg Ser Gln Val Leu Gly Arg Glu Met Arg Asp Met
65 70 75 80

Glu Gly Asp Leu Gln Glu Leu His Gln Ser Asn Thr Gly Asp Lys Ser

85

90

95

Gly Phe Gly Phe Arg Arg Gln Gly Glu Asp Asn Thr
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<210> 794

<211> 970

<212> DNA

<213> Homo sapiens

<400> 794

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<210> 795

<211> 152

<212> PRT

<213> Homo sapiens

<400> 795

Arg Pro Lys Glu Glu Val Pro Arg Ser Lys Ala Leu Glu Val Thr Lys
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Leu Ala Ile Glu Ala Gly Phe Arg His Ile Asp Ser Ala His Leu Tyr
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Asn Asn Glu Glu Gln Val Gly Leu Ala Ile Arg Ser Lys Ile Ala Asp
 35 40 45

Gly Ser Val Lys Arg Glu Asp Ile Phe Tyr Thr Ser Lys Leu Trp Ser
 50 55 60

Thr Phe His Arg Pro Glu Leu Val Arg Pro Ala Leu Glu Asn Ser Leu
 65 70 75 80

Lys Lys Ala Gln Leu Asp Tyr Val Asp Leu Tyr Leu Ile His Ser Pro
 85 90 95

006230" E6T330

Met Ser Leu Lys Pro Gly Glu Glu Leu Ser Pro Thr Asp Glu Asn Gly
 100 105 110

Lys Val Ile Phe Asp Ile Val Asp Leu Cys Thr Thr Trp Glu Ala Met
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Phe Asn Pro Gln Ala Ala Gly Asp
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<210> 796

<211> 2435

<212> DNA

<213> Homo sapiens

<400> 796

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<212> PRT

<213> Homo sapiens

<400> 797

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```

```

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```

```

Arg Gly Gly Val Gly Gly Glu Thr Arg Ala Ala Leu Ala Arg Ala Pro
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```

```

Pro Pro Gly Arg Ala Glu Trp Tyr Gly Pro Ala Gly Val Lys Ala Gly
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```

```

Gly Arg Arg Arg Val Pro Arg Arg Arg Arg Arg Trp Gly Cys Val Gln
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```

```

Glu Glu Arg Trp Ala Gly Pro Ala Arg Val Gly Gly Arg Pro Arg Gly
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```

```

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```

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<210> 798

<211> 164

<212> PRT

<213> Homo sapiens

<400> 798

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<213> Homo sapiens
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 <213> Homo sapien

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<211> 3115

<212> DNA

<213> Homo sapien

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 <211> 1238
 <212> DNA
 <213> Homo sapien

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<400> 804

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<212> PRT
<213> Homo sapiens
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Arg Leu Met Asn Arg Asp Glu Asn Gly Gly Gly Ala Gly Gly Ser Gly
          20                      25                      30

Ser His Gly Thr Leu Gly Leu Pro Ser Gly Gly Lys Cys Leu Leu Leu
          35                      40                      45

Asp Cys Arg Pro Phe Leu Ala His Ser Ala Gly Tyr Ile Leu Gly Ser
          50                      55                      60

Val Asn Val Arg Cys Asn Thr Ile Val Arg Arg Arg Ala Lys Gly Ser
          65                      70                      75                      80

Val Ser Leu Glu Gln Ile Leu Pro Ala Glu Glu Glu Val Arg Ala Arg

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85										90					95				
Leu	Arg	Ser	Gly	Leu	Tyr	Ser	Ala	Val	Ile	Val	Tyr	Asp	Glu	Arg	Ser				
			100				105						110						
Pro	Arg	Ala	Glu	Ser	Leu	Arg	Glu	Asp	Ser	Thr	Val	Ser	Leu	Val	Val				
			115				120						125						
Gln	Ala	Leu	Arg	Arg	Asn	Ala	Glu	Arg	Thr	Asp	Ile	Cys	Leu	Leu	Lys				
			130				135						140						
Gly	Gly	Tyr	Glu	Arg	Phe	Ser	Ser	Glu	Tyr	Pro	Glu	Phe	Cys	Ser	Lys				
			145				150						155			160			
Thr	Lys	Ala	Leu	Ala	Ala	Ile	Pro	Pro	Pro	Val	Pro	Pro	Ser	Ala	Thr				
				165						170						175			
Glu	Pro	Leu	Asp	Leu	Asp	Cys	Ser	Ser	Cys	Gly	Thr	Pro	Leu	His	Asp				
			180							185						190			
Gln	Glu	Gly	Pro	Val	Glu	Ile	Leu	Pro	Phe	Leu	Tyr	Leu	Gly	Ser	Ala				
			195				200						205						
Tyr	His	Ala	Ala	Arg	Arg	Asp	Met	Leu	Asp	Ala	Leu	Gly	Ile	Thr	Ala				
			210				215						220						
Leu	Leu	Asn	Val	Ser	Ser	Asp	Cys	Pro	Asn	His	Phe	Glu	Gly	His	Tyr				
			225				230						235			240			
Gln	Tyr	Lys	Cys	Ile	Pro	Val	Glu	Asp	Asn	His	Lys	Ala	Asp	Ile	Ser				
				245						250						255			
Ser	Trp	Phe	Met	Glu	Ala	Ile	Glu	Tyr	Ile	Asp	Ala	Val	Lys	Asp	Cys				
			260							265						270			
Arg	Gly	Arg	Val	Leu	Val	His	Cys	Gln	Ala	Gly	Ile	Ser	Arg	Ser	Ala				
			275				280						285						
Thr	Ile	Cys	Leu	Ala	Tyr	Leu	Met	Met	Lys	Lys	Arg	Val	Arg	Leu	Glu				
			290				295						300						
Glu	Ala	Phe	Glu	Phe	Val	Lys	Gln	Arg	Arg	Ser	Ile	Ile	Ser	Pro	Asn				
			305				310						315			320			
Phe	Ser	Phe	Met	Gly	Gln	Leu	Leu	Gln	Phe	Glu	Ser	Gln	Val	Leu	Ala				
				325						330						335			
Thr	Ser	Cys	Ala	Ala	Glu	Ala	Ala	Ser	Pro	Ser	Gly	Pro	Leu	Gly	Glu				
			340							345						350			
Arg	Gly	Lys	Thr	Pro	Ala	Thr	Pro	Thr	Ser	Gln	Phe	Val	Phe	Ser	Phe				
			355				360						365						
Pro	Val	Ser	Val	Gly	Val	His	Ser	Ala	Pro	Ser	Ser	Leu	Pro	Tyr	Leu				

370

375

380

His Ser Pro Ile Thr Thr Ser Pro Ser Cys

385

390

<210> 806

<211> 302

<212> PRT

<213> Homo sapiens

<400> 806

Val Arg Ala Arg Leu Arg Ser Gly Leu Tyr Ser Ala Val Ile Val Tyr
 5 10 15

Asp Glu Arg Ser Pro Arg Ala Glu Ser Leu Arg Glu Asp Ser Thr Val
 20 25 30

Ser Leu Val Val Gln Ala Leu Arg Arg Asn Ala Glu Arg Thr Asp Ile
 35 40 45

Cys Leu Leu Lys Gly Gly Tyr Glu Arg Phe Ser Ser Glu Tyr Pro Glu
 50 55 60

Phe Cys Ser Lys Thr Lys Ala Leu Ala Ala Ile Pro Pro Pro Val Pro
 65 70 75 80

Pro Ser Ala Thr Glu Pro Leu Asp Leu Gly Cys Ser Ser Cys Gly Thr
 85 90 95

Pro Leu His Asp Gln Gly Gly Pro Val Glu Ile Leu Pro Phe Leu Tyr
 100 105 110

Leu Gly Ser Ala Tyr His Ala Ala Arg Arg Asp Met Leu Asp Ala Leu
 115 120 125

Gly Ile Thr Ala Leu Leu Asn Val Ser Ser Asp Cys Pro Asn His Phe
 130 135 140

Glu Gly His Tyr Gln Tyr Lys Cys Ile Pro Val Glu Asp Asn His Lys
 145 150 155 160

Ala Asp Ile Ser Ser Trp Phe Met Glu Ala Ile Glu Tyr Ile Asp Ala
 165 170 175

Val Lys Asp Cys Arg Gly Arg Val Leu Val His Cys Gln Ala Gly Ile
 180 185 190

Ser Arg Ser Ala Thr Ile Cys Leu Ala Tyr Leu Met Met Lys Lys Arg
 195 200 205

Val Arg Leu Glu Glu Ala Phe Glu Phe Val Lys Gln Arg Arg Ser Ile
 210 215 220

006230" E.96T5960

Ile Ser Pro Asn Phe Ser Phe Met Gly Gln Leu Leu Gln Phe Glu Ser
 225 230 235 240

Gln Val Leu Ala Thr Ser Cys Ala Ala Glu Ala Ala Ser Pro Ser Gly
 245 250 255

Pro Leu Arg Glu Arg Gly Lys Thr Pro Ala Thr Pro Thr Ser Gln Phe
 260 265 270

Val Phe Ser Phe Pro Val Ser Val Gly Val His Ser Ala Pro Ser Ser
 275 280 285

Leu Pro Tyr Leu His Ser Pro Ile Thr Thr Ser Pro Ser Cys
 290 295 300

<210> 807
 <211> 3829
 <212> DNA
 <213> Homo sapiens

<400> 807
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 ccaagtgaag acatgcttac ttccccttca ccttccttca tgatgtggga agagtgcctgc 120
 aacccagccc tagccaacgc cgcatgagag ggagtgtgcc gagggcttct gagaagggtt 180
 ctctcacatc tagaaagaag cgcttaagat gtggcagccc ctcttcttca agtggctctt 240
 gtectgttgc cctgggagtt ctcaaattgc tgcagcagcc tccacccagc ctgaggatga 300
 catcaatata cagaggaaga agagtcagga aaagatgaga gaagttacag actctcctgg 360
 gcgaccccgga gagcttacca ttctcagac ttcttcacat ggtgctaaca gatttgctcc 420
 taaaagtaaa gctctagagg ccgtcaaatt ggcaatagaa gccgggttcc accatattga 480
 ttctgcacat gtttacaata atgaggagca ggttgactg gccatccgaa gcaagattgc 540
 agatggcagt gtgaagagag aagacatatt ctacacttca aagctttgga gcaattccca 600
 tcgaccagag ttggtccgac cagccttgga aaggctactg aaaaatcttc aattggacta 660
 tgttgacctc tatcttattc attttccagt gtctgtaaag ccagggtgagg aagtgatccc 720
 aaaagatgaa aatggaaaaa tactatttga cacagtggat ctctgtgcca catgggaggc 780
 catggagaag tgtaaagatg caggattggc caagtccatc ggggtgtcca acttcaacca 840
 caggctgctg gagatgatcc tcaacaagcc agggctcaag tacaagcctg tctgcaacca 900
 ggtggaatgt catccttact tcaaccagag aaaactgctg gatttctgca agtcaaaaaga 960
 cattgttctg gttgcctata gtgctctggg atcccatcga gaagaaccat ggttggaacc 1020
 gaactccccg gtgctcttgg aggacccagt cctttgtgcc ttggcaaaaa agcacaagcg 1080
 aacccagccc ctgattgccc tgcgctacca gctgcagcgt ggggttgtgg tcctggccaa 1140
 gagctacaat gagcagcgca tcagacagaa cgtgcagggtg tttgaattcc agttgacttc 1200
 agaggagatg aaagccatag atggcctaaa cagaaatgtg cgatatttga ccttgatat 1260
 ttttgctggc ccccttaatt atccattttc tgatgaatat taacatggag ggcattgcat 1320
 gaggtctgcc agaaggccct gcgtgtggat ggtgacacag aggatggctc tatgctggtg 1380
 actggacaca tcgcctctgg ttaaactctc cctgcttggc gacttcagta agctacagct 1440
 aagcccatcg gccggaaaag aaagacaata attttgtttt tcattttgaa aaaattaaat 1500
 gctctctcct aaagattcct cactactttt gctctccata acttctatgt tttctctcct 1560
 tctgacacac tagtgcccc aaattgtgat ttgcctatac gtttagggcc gggattggaa 1620
 gatgtaaca accatttaag attcatttct gcagtgggag tgggtggagt ttcacctct 1680
 gggaaagggg caggtgacag gtatttatca gtcagtgcct ctctagctct tgtaggaaga 1740
 agcacacgca ggatggagtc tagaggatga gcgatattga ccagcaattc atgggctccc 1800
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 tcattatatac ttgttcataa aagaaaactt tccacattgt ttttaacaaac cccacagctg 2040
 agagtcaggc ctgaatcttt gatgtgtgcc cattcacaac gttgacccta ttggtttgtg 2100
 gtggggcagg acatcgaaga tatcattgac taatcacatt cccctgaata gctcatatct 2160
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 gttttatact ttaaacaagg catagttaca agtataaaac ataaatatcc caaagccatt 2280
 atgcatggca ctcaagatta aaatgggaaa taatacatct aataaatcaa atgttccaag 2340
 acttcaaatg tcttttgga acaggctatg taaaacagca cactggtttc aaactttggg 2400
 aaattttaag aagaactctt acaaaggcat ttaattctta tacataattt tcaggggacc 2460
 taagttaatc agctaatac gaagacatga ttttcgtttt agaaaacact tttgaaaact 2520
 tgggataatc tcatgtctta atgatcaaag cattatgaga aggacagtgg ttttttacct 2580
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 catcagagat tccttctcaa gccatgtctc agagctgata ggcattcccag caagttttgc 2700
 agtcacaaat ttttctgtaa attacttatt ctataaaatt ggaagaggcc ataaactttg 2760
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 ggccatcctg ttctacctta actttctgag tctatggaat gataatttca catctcataa 3060
 acttgactga tgtaagtgtc aagaaaagat tgacattttg ttaaaacttc gtagccaagt 3120
 gtgtaacgct taagcagact ttcatatttc aaatctctat agcacgtgta actctttttt 3180
 caagatgtga aataatcatt aggtcagtc tttgtaaata gtacagctgc tgtgggcttt 3240
 ttccagttct tcaccatcca tttttataaa actcttattg ttaaaaaaaa aaagttactc 3300
 agaatttcat aaagccaaac acctgatttc aggaacactt gagatgtaag aaaattttat 3360
 agggacctcc aatcactaat tttctattt tttctctcaa agaaatgctg aaggagggaa 3420
 ttcaggttga atgaaaggaa atagtaactt acagccatat agagttataa agacttcttg 3480
 taaatgtgaa catatggtaa aatataaaaa catgtatttt tgaaaaaatg gattctactc 3540
 attattttac ttccatttaa gatataaatg tagagaaata agtataattc taagctaata 3600
 cgtacgcaat gtaggaagct gtaattactg accaaaacta tgtgaagtgg agaaaacctg 3660
 gggaagtgga tggttttaga tgaaactgaa gttaaattca tattgattta aagtaaattg 3720
 ttataacttt ataaagtttt tcatcatcac cacagcaatc acaaagagaa taattatgaa 3780
 tatacgcaag aggaaatgag aagggaatcc aaatgtcatt aaaaaaaaaa 3829

<210> 808

<211> 781

<212> DNA

<213> Homo sapiens

<400> 808

gcggcggagc tgtgagccgg cgactcgggt ccttgaggtc tggattcttt ctccgctact 60
 gagacacggc gggtaggtcc acaggcagat ccaactggga gttgaagtgt gaggtagagt 120
 gaagaggaac cagcaggctt ccggagggtt gtgtggtcag tgactcagag tgagaaggcc 180
 ctccaagtgc tgcctcctct catgcggtgc cagccccatg gaccttcttg tctcgtcacg 240
 gccataacta gggaggaagg agggccgagg agtggagggg ctgaggcgaa gctgggggtgc 300
 tgttgggggt atccgagtcc cagaagcacc tggaaacccg acagaagatt ctggactccc 360
 cagacgggac caggagaggg acggcatgag cgacacacac aaacacagaa ccacacagcc 420
 agtcccagga gccagtaat ggagagcccc aaaaagaaga accagcagct gaaagtccggg 480
 atcctacacc tgggcagcag acagaagaag atcaggatac agctgagatc ccagtgcgcg 540
 acatggaagg tgatctgcaa gagctgcac agtcaaacac cggggataaa tctggatttg 600
 ggttccggcg tcaagggtgaa gataatacct aaagaggaac actgtaaaat gccagaagca 660
 ggtgaagagc aaccacaagt ttaaatgaag acaagctgaa acaacgcaag ctggttttat 720
 attagatatt tgacttaaac tatctcaata aagttttgca gctttcacca aaaaaaaaaa 780

781

```
<400> 809
Met Arg Cys His Ala His Gly Pro Ser Cys Leu Val Thr Ala Ile Thr
          5              10              15
```

Cys Cys Trp Gly Tyr Pro Ser Pro Arg Ser Thr Trp Asn Pro Asp Arg
35 40 45

His	Thr	Gln	Thr	Gln	Asn	His	Thr	Ala	Ser	Pro	Arg	Ser	Pro	Val	Met
65					70					75					80

Glu Ser Pro Lys Lys Lys Asn Gln Gln Leu Lys Val Gly Ile Leu His
85 90 95

Leu Gly Ser Arg Gln Lys Lys Ile Arg Ile Gln Leu Arg Ser Gln Cys
100 105 110

Ala Thr Trp Lys Val Ile Cys Lys Ser Cys Ile Ser Gln Thr Pro Gly
115 120 125

Ile Asn Leu Asp Leu Gly Ser Gly Val Lys Val Lys Ile Ile Pro Lys
130 135 140

Glu Glu His Cys Lys Met Pro Glu Ala Gly Glu Glu Gln Pro Gln Val
145 150 155 160

```
<210> 810
<211> 624
<212> DNA
<213> Homo sapiens
```

```
<220>  
<221> misc_feature  
<222> (1)...(624)  
<223> n=A,T,C or G
```

```
<400> 810
atganaagga gatgacaca aagttagatc tcatcacaag tgatttggca gattaccagc 60
agccctcat gatnggcacc gggacagtca cgaggaaggg ctccaccttc cggccatgg 120
acacggatgc cgaggaggca ggggtgagca ccgatgccgg cggccactat gactgcccg 180
```

```

agcggggccgg ccgccacgag tacgcgctgc ccttggcgcc cccggagccc gactacgcca 240
cgcccatcgt ggagcggcac gtgctgcgcg cccacacgtt ctctgcgcag agcggctacc 300
gcgtcccagg gcccagccc ggccacaaac actccctctc ctcgggcggc ttctcccccg 360
tagcgggtgt gggcgcccag gacggagact atcaaaggcc acacagcgca cagcctgcgg 420
acaggggcta cgaccggccc aaagctgtca gcgccctcgc caccgaaagc ggacaccctg 480
actctcagaa gcccccaacg catcccggga caagtgcag ctattctgcc cccagagact 540
gcctcacacc cctcaaccag acggccatga ctgccctttt gtgaacacaa tgtgaaagaa 600
gcctgctgtg gtactgagcg tcgg                                     624

```

<210> 811

<211> 572

<212> DNA

<213> Homo sapiens

<400> 811

```

agcgggctgt gaggacgtc tgggccaggc tgcagcgca gcgttccgag ctgctgggct 60
ctttcgagga tgttctgata cgcgcgtcgg cctgcctgga ggaggcggcc cgggagcgcg 120
acggcctgga gcaggcgctg cggaggcgcg agagcgagca cgagaggag gtgcgcgctc 180
tgtacgagga gacggagcag cttcgggagc agagccggcg ccgcgcgagt cagaacttcg 240
cccgcgggga gcggagaagc cgtctggagc tggagctgca gatccgcgag caggaccttg 300
aacgcgcggg cctgcggcag cgggagttag agcagcagct gcacgcccag gctgcggagc 360
acctggaggc acaggcccag aactcccagc tgtggcgggc gcacgaggcg ctgcgaacgc 420
agctggaggg ggcgaggag cagatccgca ggctggagag cgaagcacga ggccgcccagg 480
agcaaaccce acgagacgtg gtcccgctct ccaggaacat gcagaaagag aaagtcagcc 540
tgctacggca actggagctg ctcagggagc tg                                     572

```

<210> 812

<211> 594

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (1)...(594)

<223> n=A,T,C or G

<400> 812

```

cggaagttag cgcagcgcg ttgccaatgg tcgctccctg atttnatgcc gctcgtggtg 60
ttttgcgggc tgccgtacag cggcaagagc cggcgtgctg aagagttgcg cgtggcgctg 120
gctgccgagg gccgcgcgg gtacgtggtg gacgacgcag ctgtcctggg cgcagaggac 180
ccagcggtgt acggcgattc tgcccgtgag aaggcattgc gtggagctct gcgagcctcc 240
gtggaacggc gctgagtcg ccacgacgtg gtcacctgg actcgcttaa ctacatcaaa 300
ggtttccggt acgagctcta ctgcctggca cgggcggcgc gcaccccgct ctgcctggtc 360
tactgcgtac ggcccggcgg cccgatcgcg ggacctcagg tggcgggcgc gaacgagaac 420
cctggccgga acgtcagtg gagttggcgg ccacgcgctg aggaggacgg gagagcccag 480
gcggcgggca gcagcgtcct cagggaaact catactgcgg actctgtagt aaatggaagt 540
gcccaggccg acgtacccaa ggaactggag cgagaagaat ccggggctgc ggag                                     594

```

<210> 813

<211> 561

ggcacgagat	ataatcagac	tcttactcct	gtacttcttag	aaatgatgca	aacacttcaa	60
ggacccacaa	atgtggaaga	tatgaatgca	ctgttaatca	aagatgctgt	gtataatgct	120
gttgatttaa	gctgccttat	agctccttta	caagtgttat	tttgatcagt	ggtttaaaaa	180
ccagcttctt	ccagaattac	aagtcattca	caataggtat	aagccaattgc	gacgcagggt	240
cattttgqctc	atcggtcagt	ggattttctgt	gaaattccaag	tctgacttaa	gacccatgct	300

```

ttatgaagca atctgtaact tgcttcaaga tcaagattta gtggccgtat tgaaacagct 360
acaactttga agttaactgt tgatgatttt gaatttagaa cagatcagtt tctaccgtat 420
ttggaaacca tgttcacact actttttcag ttactgcagc aagttacaga atgtgacaca 480
aagatgcatg ttttgcatgt cctttcttgt gtgatcgaaa gagtcaacat gcagatacga 540
ccatatgtgg gatgtttggt acaatatattg cccctccttt ggaagcagaa gtgaanaaca 600
caatatgttg agatgtgcta ttttgaccac acttattcat cttggtcagg gattangagc 660
agacagcaag acctgtccct ttcctgctcc agttattcac tgagtaccag atgtttcaca 720
gccttcncat gtttattttt ctggaaaatg ggtaaanaat atnggtanga acctttggga 780
aaac 784

```

```

<210> 816
<211> 813
<212> DNA
<213> Homo sapiens

```

```

<220>
<221> misc_feature
<222> (1)...(813)
<223> n=A,T,C or G

```

```

<400> 816
ggcacgagca ggctgggaag aagtccttgc ttctcaaggc cacgtaccgg ccgcgtcctt 60
ccacccttgc cctttaaacc acagatgcc aatgatacgc caacagacac tacattcccc 120
agcagctgct gccagagccc tctttagct tctttatttt ctgtttcttt ccagctttcc 180
taccctccta tcccccttg tgtttgggccc acaattttga aataattttt attataggta 240
tgtgtgacca aagccagatt tttataaggt aaaataaatt aagaatttaa acagtaaaag 300
ccagtgtctc aaaatgtcag cattaanaatg tgaaggggac agcaggggtgt gaaccggaaa 360
cacacattgc caaacagttg ccaactgaac tgctgcttct catgggtcgt tcttttcttt 420
gcccttaagg tcaatgccag tgtccagacg agcagtgtag aaaagctccc tgtgtgggtt 480
gtcgtgaggt ctgcttgtat ctcttctact gcgttagttt cattagctct ttattctcct 540
tacgttcgag tgaatctgcc aagaacactg gtggatagta ttatcctaac acttttggtt 600
tgggggctgg gagggggcag ggaatagtga gctggcttta ccaccttcag gatctcgaat 660
tgggcgcttg aacctaagaa agattgtgga cttatcaaaa gtcaccgctc agtggttcgtc 720
aagcatgtat ttatgtgacn atcatactag ggaggggatg gttgggaatt cttccatgtg 780
caaatttngn cccgcaanaa gcaaaaactgg ngt 813

```

```

<210> 817
<211> 229
<212> DNA
<213> Homo sapiens

```

```

<220>
<221> misc_feature
<222> (1)...(229)
<223> n=A,T,C or G

```

```

<400> 817
gaaactttta cattaatgat ttattaaaaa aaacaactcc ttgtcccact ccactgngct 60
gcttgtaatc tccatacatg gcctccattt tcaactgttt tnttggtcac anagctccaa 120
acanacacat ttttttttcc aggtaaaagc tgtttttagt ttgtagtaca aatgtgactg 180
catccaatac tgacacattg ttcctttggc ccacagtccc antcaccac 229

```

```

<210> 818
<211> 781

```

<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (1)...(781)
<223> n=A,T,C or G

<400> 818
ggcacgaggt gtgtgtgtgt gtgtgtgtgt aacacatggg cattgggcct tccaggacaa 60
cttggttagg gctccagggt ggctctcag gcaggaacag gcttttttcc tcctgtcttt 120
tcctcacatc acgtcctgcc ccaggtcact gcataaataa gtgctttgga aagtattcat 180
ctagaaagta acataaatac tgtacataga aaagggttgc cgccccttag ccttcgcact 240
gccccagaga gctctccaca tattgcacac ggctcccca gccctgtggg gtccaggcct 300
ggctgtgtct ttggtagaag cttcagggac agttcctggg cagccccac atctncaccc 360
tgctcccaa gggagctct agggtagtca gtgggtacca gaagccttgc tcggcctcgc 420
tggtggcctt ctaccangga tgctttcaca aggatgagac agaatcccaa tggtagccc 480
ctgcttggac actctgctca aggtctgcat gtggcctggg aggagacagg caggctgang 540
gcagggtggac aggtgantcc tggccacana aggcaggctc acacccttca cangaatagg 600
tggtttgngc tgtcatctcg gccacggtc tcctnntgcg ccaccccccc ttnttgaatc 660
gnaantctc aaanccctta ccaccacttg atgaccnanc atttttangg cctggcttga 720
aggngggggc cttnggcccc ccnaaggggg aaatncccc ggngaattnc ccaangggga 780
a 781

<210> 819
<211> 199
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (1)...(199)
<223> n=A,T,C or G

<400> 819
cnnngtggaa anggctgggn nngcggccgt tttcgnngta gtatcgcnt tttttttttt 60
tttttgtggg aggttntgcn gtntttgntt gctctctcaa attccaggaa ttgacttatt 120
taattaatgc ctgcaacctg tgctagcaaa tatttgnaca aaacnanttg tgttggngat 180
gttcttttgg gtcgggcag 199

<210> 820
<211> 211
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (1)...(211)
<223> n=A,T,C or G

<400> 820
nnnggcacga ggagagagag agagagagag agagagagag agagagagag agagagagag 60
agagagagag agagagagag agagagagag agagagagag agagagagag agagagagag 120
agacagtnc ntgtgtgtct ctctgtctcn aagtaacnc tgaggatct gntntctgtn 180

tntgngtaca cngtatctct cntggncata t

211

<210> 821
<211> 952
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (1)...(952)
<223> n=A,T,C or G

<400> 821
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ggcagcagag caccaaggaa acatccagac atgcgcggcc cggcccatcc gctcccggaa 120
cagcaccaag acgaaatggg aaactacatg tccccagggt cgaggctgca ggggcagact 180
ctggtgtgaa caggggggat gtgaccacct aaggaaaagg tcacacctgt cttggtatca 240
ggggctcaag agctctcaaa aatgtaaggg gccgacagtc ccctgccccca ggctgatca 300
caactccagg gtcctgaggt cagagtaaag tgcagagggt tttaaacata accaaaattt 360
caggagaggc caattcttac ttgaaagagc aacacctggg ggcgctgctt gccattactt 420
cctcatcttt agcaacacat ttgcttttca aggtgttctt tgtggaaaca cacatacaca 480
tagacacatg cccctcagat gtccccctgcc ccctgattag tagaatgtgg gggtttccaca 540
atgagcagaa actgatccaa ttttggttaa gtttgagaag ccctctgaat ttgggtgggt 600
ggcccaatgt aaatacttcc gcagagatgg agggcattca aaacagggtc tgaaaggatc 660
cagcctatct tggactttgt tctggaancc anggattcag cnttggccac ctgtgccagg 720
cttgcaaggc ctggtgtgaa cncccaaant ggcagcaaaa acaacanaca gccnctgcac 780
tttggnntgga ccaacgtttg gcctnaacaa atctnngcggg ttgggatntt cttgntttcn 840
cncccagggg accnaaaacc ccntaentg naataacnt ttttttttnn aacntttan 900
ccantgggnt tncnaaaaa acttgncccc ttttttttnc caangnaaa at 952

<210> 822
<211> 587
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (1)...(587)
<223> n=A,T,C or G

<400> 822
ggcagcagaa ctagtctcga gttttttttt ttttttttta acatttctga attttattat 60
ttttagggaa gacacgcagt ttcacaagaa acaatgattt ttctcaaaca atagaaaaaa 120
aggtcttttt gaaaaatcca ctgtcttaga tgaaaagtct acccagcaag cactggggca 180
gttctgagag tagaaaccag tgtggtggaa gttacttata ggaagttagc tgcagagggtc 240
tcacaaagtc ctgattagtt ctgnaaggct ccattgggcc agctcagggt aacagtggga 300
atgagctcac agacaaaaggc aggcaccagt tcctntgccc gggatgcagg ctggctcact 360
ccccangcgg ntgcactctt ctccagactc atcaaactgc tgctgtccan ctncgncatg 420
actntgttga gaacatanaa ctctgctctc tggctttgct tcanctcctg gtgggcnnaa 480
ttctgcttag ccttctncac tntgaaggnt gggcttttaa ctttttgatt tttttttcn 540
ggcaggggga accatgaatg gggtagatac ccacncnggg ntttggc 587

<210> 823
<211> 264

<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (1)...(264)
<223> n=A,T,C or G

<400> 823
ntcnatncct actangncaa actgactccg ccctnagnca cctngtggtc canggctgcg 60
gagctgcgat acagccttcc ggggtctgn tggaaccccg acctntcntg gtgtntntcc 120
ntcccnccc ccaacccgcc aagggcctgc ctttccctnct gggcctttgc cagcgnnngg 180
ccanaccggg gccaaaccgg nccccgggca cattttaacc nagggcncnc ttntagaana 240
aaaccccggn tgatgttata aagg 264

<210> 824
<211> 520
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (1)...(520)
<223> n=A,T,C or G

<400> 824
tcaagcngcc cccantntga tggatatctg caaaattcnc cctttcacccg gccgcccgc 60
gcatgtctta ttatacaaca natccaactt ccctaagnng ntcacacatn ntaagggtatt 120
gttaacaaaa taggaaantc tattngaact aacaatcatc tctttgaatc tgcntatccc 180
attaaaagca ttttctctca tattcctcat atcggttatg gncaatggat acccatctga 240
gctgggtgan ccttttaaat tnattatact taactttttg aaggctgtta taccacaagg 300
acaaacctaa ncaaccanca gatatacttg anggtntctc ctgtnatttc tcagattcca 360
atataccatt ttgccttnac acctacagcc cttaggggca tctcntttcc ncanaacaaa 420
ncattntcac taagacagnc tggggtntntn caccaatggc taccaaacct ctgnccgcna 480
cccaccgcnt aaanggcnga aattncnanc ccacacgggt 520

<210> 825
<211> 2064
<212> DNA
<213> Homo sapiens

<400> 825
cggtgcgctg agcgccggag gagcgtaggc agggcagcgc tggcgccagt ggcgacagga 60
gccgcgcgac cggcaaaaaat acacgggagg ccgtcgccga aaagagtccg cggtcctctc 120
tcgtaaacac actctcctcc accggcgccct cccctccgc tctgcgcgcc gcccggtg 180
gcgcccagag cgcctccgac tgctatgtga ccgcgaggct gcgggaggaa ggggacaggg 240
aagaagaggc tctcccgcgg gagcccttga ggaccaagt tgccggccact tctgcaggcg 300
tcccttctta gctctcgccc gccctttct gacacctagg cggcccggt tctcttctct 360
tctcgcgcgc cccagcgcgc tcggttcccc gcgacctagg tgacgatgga ggagctgcgg 420
gagatggact gcagtgtgct caaaaggctg atgaaccggg acgagaatgg cggcggcgcg 480
ggcggcagcg gcagccacgg caccctgggg ctgccgagcg gcggcaagt cctgctgctg 540
gactgcagac cgttctctgg gcacagcgcg ggctacatcc taggttcggt caacgtgcgc 600
tgtaaacacca tcgtgcggcg gcgggctaag ggctccgtga gcctggagca gatcctgccc 660
gccgaggagg aggtacgcgc ccgcttgccg tccggcctct actcggcggt catcgtctac 720

Glu Pro Leu Asp Leu Gly Cys Ser Ser Cys Gly Thr Pro Leu His Asp
180 185 190


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ctgactcttc catctgtgca ggttgactga ggtcattcct gagttgcagt atgttgagag      60
ggtaatatatt ctgtctttctc taactcccca tactcccttg tcttccactc tccacttagg      120
agttttttgt gagttatgtc cttgttgctt ttgcctcttt ttctttctag ccttgattgt      180
gccagaagac aatgtcccta ttcacacact ctttctgctt ttctgtgggc aggaacatgg      240
aaggggtgct gatggacgtg gactgtgaga gcgtctaccc cactgtgtag g                291

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<210> 833
<211> 491
<212> DNA
<213> Homo sapien

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<400> 833
ctgtagcttc tgtgggactt ccactgctca ggcgtcaggc tcaggtagct gctggccgcg      60
tacttggtgt tgctttgttt ggagggtgtg gtggtctcca ctcccgcctt gacggggctg      120
ctatctgcct tccaggccac tgtcacggct tccgggtaga agtcacttat gagacacacc      180
agtgtggcct tggttgcttg aagctcctca gaggagggcg ggaacagagt gaccgagggg      240
gcagccttgg gctgacctag gacggctcagc ttggctccctc cgccgaagac cacattattg      300
ccgtcccacg tctgacagta atagtcagcc tcatccatag cctgggtccc gctgatggtc      360
agagtggctg tgttcccaga gttggagcca gagaagcgct cagggatccc tgaagaccgc      420
ttattatctt gataaatgac taccacaggg gactggcctg gcttctgttg ataccaacaa      480
gcagatacct g                491

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```

<210> 834
<211> 308
<212> DNA
<213> Homo sapien

```

```

<400> 834
ctggctgagg tccacgccgc ggtaggtgaa cttgcggaag gtccgcttct tcttctgctc      60
tacttctgcc gtgctggaga acatcgaact gaacaagaag agtatgtatt cccgtgtgcc      120
agagtgccag gtcaccacat actattatgt tgggttcgca tatttgatga tgcgtcgta      180
ccaggatgcc atccgggtct tcgccaacat cctcctctac atccagagga ccaagagcat      240
gttccagagg accacgtaca agtatgagat gattaacaag cagaatgagc agatgcatgc      300
gctgctgg                308

```

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<210> 835
<211> 472
<212> DNA
<213> Homo sapien

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<220>
<221> misc_feature
<222> (1) ... (472)
<223> n = A,T,C or G

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<400> 835
ctgacatggt aactgtgatg cataaaactc gatcttctga tggggagtaa gtgcagaagg      60
tagaaatctc cgccccgcgg gggcttatct gtactggtag ttcattgctgt ggtctgcgtt      120
tgtgccatag ccgccttgtg aggactggta ggagctggga gggccactgt agttctggcc      180
ggaccccggg gagttgtagt tcgactgtga gtagectcct tgtttgcctt ggtatgagga      240
ggcgccccca gaacctccgc cgtagcccc gtgtgacctt gggttgtagg atgccccgcc      300
tgagccgtag ctgttccgcg cgcttcggcc tccactacca ctgtagtga atttgctctc      360
gtagntgtag tcggatccgc ccccgcccc gggagagttg tngganttcg agtaggagta      420
gctgccttgt ccatggttat agcctttctg cttgcctgtt ggagggccat ag                472

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<210> 836
 <211> 354
 <212> DNA
 <213> Homo sapien

<400> 836
 ccagtgcac cttcagatag acacatgggtg accagagccc gccaggcttc tgcagggtggc 60
 agtgtcgagc aagtgtgaaga tgtctgtggg aaggagaagc tcctgaaatg aacgttctgc 120
 aaacagaagg ctgaggggtc ttccaggcat gtccagtcac taggagctgc caccgggtggg 180
 cttgagtgcc aggctctagg ctttgtgcag aaagcaccgc gggcgggggg cggttaaggga 240
 gagcaaaatg ggtctctctc aactgcagtc agtgcctctg ggaacacggc ctcacagaca 300
 gcacatattc tacgtcacag ctctagggtt tcaaggactt agccatccga cagg 354

<210> 837
 <211> 318
 <212> DNA
 <213> Homo sapien

<220>
 <221> misc_feature
 <222> (1)...(318)
 <223> n = A,T,C or G

<400> 837
 ctgaaaatga aggtaattaa aaccatggag gcgatcagcg aggttctcca ggacotttagg 60
 tttgatgcgg aatctgccga gtgatggcgg ctccccaggg atgcgccgag ggagatggga 120
 aacggggcgg atggcgccca gccagccct aactgccagc cacattgaag cggacattgg 180
 caaccgggtc cccagccatg cgcagaaccg tgggtagcat gtgcttggtg gtgatgtcct 240
 gccacagac ctcagacggc acattgatgc agaagagcgt antcatgcgg tgcaggtagt 300
 tggggtctcc ggacatgg 318

<210> 838
 <211> 277
 <212> DNA
 <213> Homo sapien

<400> 838
 ctgcgcgtcg ccaaagtgc aggcgggtgc gcctccaagc tctctaagat ccgagtcgtc 60
 cggaaatcca ttgcccgtgt tctcacagtt attaaccaga ctcagaaaga aaacctcagg 120
 aaattctaca agggcaagaa gtacaagccc ctggacctgc ggcctaagaa ggcacgtgcc 180
 atgcgcgcgc ggctcaacaa gcacgaggag aacctgaaga ccaagaagca gcagcggaag 240
 gagcggctgt acccgtctgc gaagtacgc gtcaagg 277

<210> 839
 <211> 276
 <212> DNA
 <213> Homo sapien

<400> 839
 ccaaggaatg caggctgtac tatctgcgaa atggagaacg tatttcagtg tcggcagcct 60
 ccaagctgct gtccaacatg atgtgccagt accggggcat gggcctctct atgggcagta 120
 tgatctgtgg ctgggataag aagggtcctg gactctacta cgtggatgaa catgggactc 180
 ggctctcagg aaatatgttc tccacgggta gtgggaacac ttatgcctac ggggtcatgg 240

ctgtctccttg	gcctggettt	acctgcactt	gcggcttctc	tggggtgcta	tcttgcactc	300
agccctgatc	tactttctgg	gaacctttct	gctatccata	ttgatcgctt	ggactgtgca	360
gtatttccag	tctgtctcag	caagcgatcc	ccctccaaga	ccatcccagg		410

<210> 848
 <211> 557
 <212> DNA
 <213> Homo sapien

<220>
 <221> misc_feature
 <222> (1)...(557)
 <223> n = A,T,C or G

<400> 848						
cacgggcccc	cagccctgtg	tcggccttgt	ctgtctcagc	tcaaccacag	tctgacacca	60
gagcccaactt	ccatcctctc	tggtgtgagg	cacagcgagg	gcagcatctg	gaggagctct	120
gcagcctcca	cacctaccac	gacctcccag	ggctgggctc	aggaaaaacc	agccactgct	180
ttacaggaca	gggggttgaa	gctgagcccc	gcctcacacc	cacccccatg	cactcaaaga	240
ttggatttta	cagctacttg	caattcaaaa	ttcagaagaa	taaaaaatgg	gaacatacag	300
aactctaaaa	gatagacatc	agaaattggt	aagttaagct	ttttcaaaaa	accagcaatt	360
ccccagcgta	gtcaaggggtg	gacactgcac	gctctggcat	gatgggatgg	cgaccgggca	420
agctttcttc	ctcgagatgc	tctgctgctt	gagagctatt	gctttgttaa	gatataaaaa	480
ggggtttctt	tttgtctttc	tgtaaggngg	acttcacgct	tttgattgaa	agtcctaggg	540
tgattctatt	tctgctg					557

<210> 849
 <211> 525
 <212> DNA
 <213> Homo sapien

<400> 849						
ctgatggttt	ggaaatgaga	gaactacagt	ggtgaagaga	ccaggaggca	gctctcagtg	60
aaaccaacat	tcgggatgcc	cttcgtgagc	cttctcagtc	ccagcaggaa	gcccacaaca	120
ctggcctccc	cagcctgcct	gctgacaaca	cctaggctta	ctttatctaa	aatcagagtg	180
taccaggtct	gtagcagaaa	ataatcaact	aaatgtcagg	gacctatgag	tcatttaaaa	240
caaaagagga	agtgaagcc	attaggcaag	ctatgtgctg	ggctgctaac	gtagcccttg	300
cagggagggg	tcaggagcgc	gctgcagtga	gccttgggtc	tcgcaggccc	agccctgctg	360
caaggagcca	gggcacccag	gaaacatcag	cacacacaca	cacagggacc	ctcccttcat	420
gtcacttggt	ttgctgccct	aaatggcttc	ttgcacccta	acccctgatc	ctggaagaag	480
gcagagagac	tggcccgtac	agagacctgc	aattctacgc	aagct		525

<210> 850
 <211> 384
 <212> DNA
 <213> Homo sapien

<400> 850						
cctcttggag	cacatccttt	actgcattgt	ggacagcgag	tgtaagtcaa	gggatgtgct	60
ccagagttac	tttgacctcc	tgggggagct	gatgaagttc	aacgttgatg	cattcaagag	120
attcaataaa	tatatcaaca	ccgatgcaaa	gttcaggta	ttcctgaagc	agatcaacag	180
ctccctgggtg	gactccaaca	tgctgggtgcg	ctgtgtcact	ctgtccctgg	accgatttga	240
aaaccaggtg	gatatgaaag	ttgccgaggt	actgtctgaa	tgccgcctgc	tcgcctacat	300
atcccaggtg	cccacgcaga	tgctccttct	cttccgcctc	atcaacatca	tccacgtgca	360

gacgctgacc caggagaacg tcag

384

<210> 851

<211> 423

<212> DNA

<213> Homo sapien

<400> 851

ctcaggaaaa	accagccact	gctttacagg	acaggggggtt	gaagctgagc	cccgcctcac	60
acccaccccc	atgcactcaa	agattggatt	ttacagctac	ttgcaattca	aaattcagaa	120
gaataaaaaa	tgggaacata	cagaactcta	aaagatagac	atcagaaatt	gttaagttaa	180
gcttttttcaa	aagatcagca	attccccagc	gtagtcaagg	gtggacactg	cacgctctgg	240
catgatggga	tggcgaccgg	gcaagctttc	ttcctcgaga	tgctctgctg	cttgagagct	300
attgctttgt	taagatataa	aaaggggttt	ctttttgtcc	ttctgtaagg	tggacttcca	360
gcttttgatt	gaaagtccta	gggtgattct	atttctgctg	tgatttatct	gctgaaagct	420
cag						423

<210> 852

<211> 413

<212> DNA

<213> Homo sapien

<400> 852

ctgaaaacag	tgggaggcca	gatgctggca	tcttccagac	gggagcatag	ccatggtcac	60
tctagccgat	gtctcctggg	gctctcaggc	ggcaaggacc	agatgcacca	ctactgtcca	120
atcccagttt	tacttagagc	cacctccttt	tttggggcca	ttagtcctta	tttcatgccca	180
gatttttact	agcggctccc	tgttcttcca	aatcaattca	tgaccgtaag	taacatacca	240
tattccaaaa	agagctcccc	caagatgtgc	cgcgatgatca	aaaaatttcc	atcccaggat	300
cattcctgct	gtatccatgg	cgataatggc	tttcagggcca	ttccctgctg	tgaacgtgaa	360
catcggaagg	aaaataatgg	caagcctccc	ttctggggatc	ttagtgccaga	cag	413

<210> 853

<211> 288

<212> DNA

<213> Homo sapien

<400> 853

atctgtgagt	tctgagaggc	atttaggccca	tgggacaggg	aggatcctgt	ctggccttca	60
gtttccatcc	ccaggatcca	cttggtctgt	gagatgctag	aactcccttt	caacagaatt	120
cacttggtggc	tattagagct	ggaggcacc	ttagccactt	cattcccttg	atggggcctg	180
actcttcccc	ataatcactg	accagccttg	acactcccct	tgcaaaccat	cccagcactg	240
caccccaggc	agccactcct	agccttggcc	tttggcatga	gatggggg		288

<210> 854

<211> 427

<212> DNA

<213> Homo sapien

<400> 854

ccaagtgaga	tcagccctca	agggcacatg	ccaagggcag	agcagcccat	gtagacagct	60
tcggagggga	tgggggtgta	gggagttcgg	ggtagctcct	cattaactat	ttgttgggtg	120
agtaaaaggg	tgaggctcag	tggcaggtac	ctctgcaatg	acaagctgcc	tcccctctat	180
gtgttttagca	tatgttatta	gaacgtgtcc	gacacccta	ccgctgccat	ttgggcccctt	240
taataaagcc	aagtagagaa	atctggcaat	aaaaggcaaa	tgtaagcatg	ctttctttaa	300

gacgcatcat aaatgggtttt ctttaagtga atggaagagt ttgacagaga tacacctttg 360
 taagaaaaca ttaagaatgc tggctgactg tgggtggctca cacctgtatt cccagcactt 420
 tgggagg 427

<210> 855
 <211> 311
 <212> DNA
 <213> Homo sapien

<400> 855
 ccagtattcc tggaggatat aacactgaca tcagcagggg tttcaatggc aacaattgca 60
 cgagctgccg gcagaagctt ctcccagggt ctcttgagat ttatgatata gatgccatca 120
 cttttccttt tatagatgta ctgttccatc tggaaagtcaa gattgggtgcc acctaaagtgg 180
 gttcctgctg caaggaactt aaggacatcc tcctccttca tttgcaggac atcaagggct 240
 ccggacattg tgaaagtttc cctttaagtt acgacgggaa tccagaacaa cgccgtatgg 300
 acccctctgc a 311

<210> 856
 <211> 328
 <212> DNA
 <213> Homo sapien

<400> 856
 cctatggaag tttgggtgctt tgctccctgt gtttgcgaaa caggtatctc gtgatttcag 60
 aaaagcttga ggagattaag tctttccggg agctgacctg cctggatctt tcctgttgca 120
 agcttgagaga tgagcatgaa cttctagaac atctcaccaa tgaagccctg tctagtgtaa 180
 ctgagctcca cctgaaggat aattgtctat ctgatgctgg ggtgcggaag atgacagcac 240
 cagttcgagt gatgaaaaga ggtatccaat gcctgcatct gtgatctcag ggttacatga 300
 taagtctaata aatgttagat tctcaagg 328

<210> 857
 <211> 502
 <212> DNA
 <213> Homo sapien

<220>
 <221> misc_feature
 <222> (1)...(502)
 <223> n = A,T,C or G

<400> 857
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 actgaggaag agaagaattt caaagccttc gctagtctcc gtatggcccg tgccaacgcc 120
 cggctcttcg gcatacgggc aaaaagagcc aaggaagccg cagaacagga tgttgaaaag 180
 aaaaaataaa gccctcctgg ggacttgga tcaagtcggc gtcattgctgg gtctccacgt 240
 ggtgtgtttc gtgggaacaa ctgggcctgg gatggggctt cactgctgtg acttcctcct 300
 gccaggggat ttggggcttt cttgaaagac agtccaagcc ctggataatg ctttaactttc 360
 tgtgttgaag cactgtttgg tgtttggtta gtgactgatg taaaacgggt ttcttgtggg 420
 gaggttacag aggttgactt cagagtggac ttgtgttttt tcttttttaa gangtaaggt 480
 tgggctgggtg ctcacagacc tc 502

<210> 858
 <211> 411
 <212> DNA

<210> 862

<211> 561
 <212> DNA
 <213> Homo sapien

<400> 862
 ccaggtcatc accattggca atgagcgggt ccggtgtccg gaggcgctgt tccagccttc 60
 cttcctgggt atggaatctt gcggcatcca cgagaccacc ttcaactcca tcatgaagtg 120
 tgacgtggac atccgcaaag acctgtacgc caacacgggt ctgtcgggcg gcaccaccat 180
 gtatccgggc attgccgaca ggatgcagaa ggagatcacc gccctggcgc ccagcaccat 240
 gaagatcaag atcatcgcac cccagagcgc caagtactcg gtgtggatcg gtggctccat 300
 cctggcctca ctgtccacct tccagcagat gtggattagc aagcaggagt acgacgagtc 360
 gggcccctcc atcgtccacc gcaaattgctt ctaaaccggac tcagcagatg cgtagcattt 420
 gctgcatggg ttaattgaga atagaaattt gccctgggca aatgcacaca cctcatgcta 480
 gcctcacgaa actggaataa gccctcgaaa agaaattgtc cttgaagctt gtatctgata 540
 tcagcactgg attgtagaac t 561

<210> 863
 <211> 291
 <212> DNA
 <213> Homo sapien

<400> 863
 ccatagctgt cccacctatg gttttaaaaa cagactgtaa cttgatcttc tgaaatcctt 60
 ctgaaccac aactcgttct gttaaagaaa tcctaggaaa gaagtcctac tgatattgtc 120
 gatagtctcc aaaagggtgag gaaggtaact gagttgaagg caactgggag gggctctctg 180
 caaactgagg accattggaa aactgtgcag aggcaaattc tgtcaacaag ataccagctc 240
 cttcaattaa agctaggaga atgccacca ttgcggctga cccaaccatg g 291

<210> 864
 <211> 265
 <212> DNA
 <213> Homo sapien

<400> 864
 ctgaactttt ccacctggag tccttgggaa taccggacgt gatcttcttt tatagggtcca 60
 atgatgtgac ccagtcctgc agttctggga gatcaaccac catccgcgtc aggtgcagtc 120
 cacagaaaac tgtccctgga ggtttgctgc tgccaggaaac gtgctcagat gggacctgtg 180
 atggctgcaa cttccacttc ctgtgggaga gcgcggctgc ttgcccgtc tgctcagtgg 240
 ctgactacca tgctatcgtc agcag 265

<210> 865
 <211> 144
 <212> DNA
 <213> Homo sapien

<400> 865
 cctccacctg cgttttgatc tagatgagca tattgtccat ctcccacagc ttgctccggt 60
 tccgcaggta cggccgcccg tgctcgcgcg tcagcgacgc gatgtcctcg cgcactcgt 120
 tgatgaccgg gagcagaaac tgct 144

<210> 866
 <211> 241
 <212> DNA
 <213> Homo sapien

<400> 866
 ctggctgtaa gtagcttcat agcaccagtc tttgagaatg tcaagctctc cagaaatcat 60
 ggcctccagg acattgggga tgatgtcggt ctgcactgt ttcagaaacc ggtccttgtc 120
 aaaggccggg tccaccggga ggatctccgt gagcacctcc gacatctctg tcttgagaa 180
 caggccccc agcaagtcgg tgacctgtc cgtaaggggc cgggatgcc ggatgaacgc 240
 g 241

<210> 867
 <211> 364
 <212> DNA
 <213> Homo sapien

<400> 867
 cctgggcccg ctgacttcag ggtgaggcca cagctactgc agcgcttttt atttatttat 60
 ttatttactg agatggagtc ttgctctgtc acccaggctg gagtgcagtg gtgcaatctc 120
 ggctcactgc aacctctgcc tcctgggctg cagtgtattct cctgcgttca agtaattctc 180
 ctgcctcggc cttctgagta gttgggatta caggcatatg ccaccacact tggctaattt 240
 ttcgatattt tagtagaaat ggggtttcac catgttggcg aggctgggtc cgaactcctg 300
 acctcaagga tcctcctgcc tcggcctcct aagggtgctgg gattgcagggt gtgagccacc 360
 acgt 364

<210> 868
 <211> 472
 <212> DNA
 <213> Homo sapien

<400> 868
 ccaccagtcc acagatgtga ctggtaaggg atctagtaac agaggatgga gttgggcaga 60
 atattatcct ggatgatatg caccagcac taggatacac ctttcattag aatgaagaga 120
 acagacaaag cctcagaaa agatacaaag gcagagacat tgattagaac attatctcat 180
 aacagagggtg gggccattac ccaccattat tgtaaaataa ctgtaactaa ccaaaacaca 240
 tacaggcttc ttaaatggag ttaataaaaac tatggcacat tgggaatcag gggcagaggt 300
 actgtttcca gacggaaaac tgggataaag ggagccatgc tgacagggcc ttattccagt 360
 ctaggttgtt agaaaggagc cctagcccag aaatgacagc aaatagccat aatcattatg 420
 tggggctgaa ccagaggaag ccaggctgag ccaagaagct ggaagtatct tg 472

<210> 869
 <211> 368
 <212> DNA
 <213> Homo sapien

<400> 869
 cttttcttgt aagtgaagaa aaaggaatgc agcaaagaag agttcgacat tggagtcctt 60
 agttccatca ggatccatt cgcagccttt agcatcatgt agaagcaaac tgcacctatg 120
 gctgagatag gtgcaatgac ctacaagatt ttgtgttttc tagctgtcca ggaaaagcca 180
 tcttcagtct tgctgacagt caaagagcaa gtgaaaccat ttccagccta aactacataa 240
 aagcagccga accaatgatt aaagacctct aaggctccat aatcatcatt aaatatgcc 300
 aaactcattg tgacttttta ttttatatac aggattaaaa tcaacattaa atcatcttat 360
 ttacatgg 368

<210> 870
 <211> 411
 <212> DNA

<400> 873
ctgtgggctc tgaatggcgt cccctttggct atccacgccg ccggcgacca ctgaattctg 60
tggttctaca acagggctctg gctgaccgaa ttgtcagaga cgtccaggaa ttcattcgata 120

```
<210> 874
<211> 156
<212> DNA
<213> Homo sapien
```

```
<210> 875
<211> 512
<212> DNA
<213> Homo sapien
```

<400>	875						
ccagcatagc	gaaaaacttgt	ctctactaaa	aatacaaaaa	ttagtcaggc	atggtggtgc		60
acgtctgtaa	taccagcttc	tcaggaggct	gaggcacgag	gatcacttga	accaggagg		120
aggaggttgc	agtgagctga	gatcatgcc	gggcaacaga	atgagacttt	gtttaaaaaa		180
aaaaaaagt	acttgattta	agggaaaaaa	tgactggcta	tattcagtca	gatatggcaa		240
agagtctcaa	ggtgttaatg	tgaatgatta	aggtcttggg	gggggtgtcc	cctatcagac		300
tacaggtgtt	tagaggcaca	gaaaaagggtg	cagttgggtt	cttaatgtga	aatgatgaga		360
agcacaactc	cagtgtgtct	ctttgtgtag	aatgtcagca	gacacccct	gctagatgtg		420
ctggatcatg	ggaagcatt	tccatttgtt	aatagattgt	tcagaagttt	taatttatga		480
tgggtgtggt	ggctcatgcc	tgtngtccca	gc				512

```
<210> 876
<211> 199
<212> DNA
<213> Homo sapien
```

<400>	876						
cctgtgccgg	gccccagggc	tggcagccac	cagctcctct	tccaggcatg	ggggacaccc		60
tgacaggatc	cggaagtctc	catttaccca	aaaatgcaag	agccatgatc	agtcatggcg		120
acactgcagc	cggtactgag	tgaccatgtc	cagtcgggct	ccgtccctcc	cacacggggg		180
acaaqcttct	ccgaggagg						199

```
<210> 877
<211> 486
<212> DNA
<213> Homo sapien
```

<400> 877

ccatcccaca taccctcact ggcattccagg agaccagcag cagggtcaag accccaaatg 360
 ttgggcacca caaataatgt gatatgtgcc aggagcacgg ggggtagggg tgaaagagaa 420
 aaacaataag g 431

<210> 881
 <211> 335
 <212> DNA
 <213> Homo sapien

<400> 881
 ccacagaggt ggtattacaa aatatacaaa gtgggtttctt tctttacatt tcatagaaga 60
 agcctgcctc atttccaaat gagagcacta gaagcacaaa tcatgcagac catttactat 120
 ataacttatg aaaaatgctg tacagggctg tgactataga tatagagtat ttggctctgt 180
 ttgggaattg atatctacaa gggggagggt caggggagga ctgtctgata tcttgacttg 240
 ctgggatggt ggagaagctg ggatggggga ggccccaatc ttgctgcacg gctacacca 300
 ctctctcttt cctagataag gctggagcgc actgg 335

<210> 882
 <211> 353
 <212> DNA
 <213> Homo sapien

<400> 882
 atgcactcaa agattggatt ttacagctac ttgcaattca aaattcagaa gaataaaaaa 60
 tgggaacata cagaactcta aaagatagac atcagaaatt gttaagttaa gctttttcaa 120
 aaaatcagca attccccagc gtagtcaagg gtggacactg cacgctctgg catgatggga 180
 tggcgaccgg gcaagctttc ttcctcgaga tgctctgctg cttgagagct attgctttgt 240
 taagatataa aaaggggttt ctttttgtct ttctgtaagg tggacttcca gcttttgatt 300
 gaaagtccta ggggtgattct atttctgctg tgatttatct gctgaaagct cag 353

<210> 883
 <211> 193
 <212> DNA
 <213> Homo sapien

<400> 883
 ctggcagaga agaatggcta cgtgactgtc agtgagatca aagccagtct taaatgggag 60
 accgagcgag cgcggaagt gccggaacac ctgctgaagg aagggttggc gtggctggac 120
 ttacaggccc caggggaggc ccactactgg ctgccagctc tctcactga cctctactcc 180
 caggagatta cag 193

<210> 884
 <211> 461
 <212> DNA
 <213> Homo sapien

<400> 884
 ctgaagaacc ccacagcgg gctgttagaa tatgccagct tcgctagtca aacctgtgag 60
 ttcaacatga tagagcagag tggaccaccc catgaacctc ggtaagagac caccaggaa 120
 ctgtacctag ggttgggggc aggtgctttt gctcctgacg cagtcttggc tgatttgtga 180
 gcagtgtgtg ttggtggcgc ctatcttttc ctcttccct tctgcctttt agctaaattc 240
 cccttgattg gccctttctc cagatattga gcagggaata tagaccttg accagccaga 300
 atcttggctg aacaaggggg aggttgactc tggtggctgt aatgaagctt ctttagaaat 360
 gattggtttt ggccgtacgc ggtggctcat gcctgtaatc ccagcacttt ttgaggccga 420

461

<211> 266

<213> Home

 $\langle 220 \rangle$ $\langle 222 \rangle \quad (1) \dots (266)$

<400> 885

<210> 886

<211> 402

<212> DNA

<400> 886

<210> 887

<211> 342

<212> DNA

<400> 887

<210> 888

<211> 228

<212> DNA

<400> 888

cgcgctcgcc	aaggctgctg	ctgttgctcc	tccaaagaag	gttggttca	aggcctgtc	60
cagggaccca	cgagcagagg	cactgggggg	caagggatct	ccaagggggc	aagggatccc	120

taaagggggt agctcacagg tgaggggggt tagggcccct ctagggagcg cctgaggcca 180
tacattcaag agtgtccctg gtgaggccca ggaagagcc aggactgg 228

<210> 889

<211> 378

<212> DNA

<213> Homo sapien

<400> 889

ttggcttttc	tcccccttc	atcctcctct	cccccttcct	caactgaaggc	tgtgagttgc	60
tttcaatgtg	acaacactat	gatgtcattt	ggaaggattt	gccaggacag	actgattctg	120
agtctctggg	gccgtatgtg	tatgcggcag	tggtgtcagg	cgatcttggt	tgaagctcta	180
tggtgccata	attaccatca	agtacacact	gttggcaaaa	ggctaacacc	tgactttagg	240
aaatgctgat	ttgagaacaa	aaggaaaggt	cttttttcac	tgcttaaagt	ggggtcactt	300
tgataccttt	gcggtcatgt	ctgtgtctga	tgagtgtaga	atctctggat	gtgcactgtc	360
agtcattgtg	ccaccagg					378

<210> 890

<211> 215

<212> DNA

<213> Homo sapien

<400> 890

ccattttgga	gtgtgtccat	tggttagcaa	tgtggaaacc	accagggcct	ttgtggagaa	60
aatggagggg	gttgaggagg	tcccaggagg	ggcttatattg	agggcctttg	ccacttgctc	120
ataggcgagc	tcgatctcct	catcatctgg	acaggtggaa	gcgaattctt	cccgggcgta	180
ggcattgctc	aagtaccgat	gcactccccg	gaagg			215

<210> 891

<211> 412

<212> DNA

<213> Homo sapien

<400> 891

ctgggtcaagt	tcaacagagc	cttggctgac	cattctatgg	ctcaggcacc	tcggctcatt	60
gatggcattg	ttcttaccac	atttgatacc	attgatgaca	aggtgggagc	tgctatttct	120
atgacgtaca	tcacaagcaa	acccatcgtc	tttgtgggca	ccggccagac	ctactgtgac	180
ctacgcagcc	tcaatgccaa	ggctgtgggt	gctgccttca	tgaaggctta	acgtggctct	240
tgcccaatac	caaatcgccg	ctttccccac	aagcccttct	tcctgtatca	agaatgtgct	300
ttagagtatg	tgagcaacct	gtcttcagtg	tagtacaaag	gcagagttag	ggggcttgtg	360
gctccttcca	acccactcc	ccgttcagca	cagccgccat	ctgcaaggaa	gg	412

<210> 892

<211> 472

<212> DNA

<213> Homo sapien

<220>

<221> misc_feature

<222> (1)...(472)

<223> n = A,T,C or G

<400> 892

tttttttttt	tttttttttt	ttaattacta	ccttttattc	taatgtgaac	catggccctg	60
------------	------------	------------	------------	------------	------------	----

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aaagctgata acaagcttgg ctgancagag ggaactaggg gtcggcagaa aggattatgg      120
gtggaaaaca ttggctcttc cttggggagt gatgctgggg aaaggggaana nagtgggtca      180
ncctgcaggt aaataggcta naaaagccaa ggccaaaggc tggaggggag aggacagtca      240
gcatgtccag cctgggggtct ggggtgtagg ttatcccttc tccctgtgcc tccccatctc      300
gtccatgagc ctagggtcttg gagccttgtg ttggaggctg ctgtgatgtc aggaacgggg      360
atctgtctag cttttggcca cttcctggga cctcacgccc ctgttgacag atggagattg      420
ggcagcaggg ccttgctgcg ttgttatctg ctgttccgac ttggtttgtc tt              472

```

<210> 893

<211> 477

<212> DNA

<213> Homo sapien

<220>

<221> misc_feature

<222> (1) ... (477)

<223> n = A,T,C or G

<400> 893

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caaagattca ctttatattat tcattctcct ccaacattag cataattaaa gccaaaggagg      60
aggagggggg tgaggtgaaa gatgagctgg aggaccgcaa taggggtagg tccccgtgg      120
aaaaagggtc agaggccaaa ggatgggagg gggtcaggct ggaactgagg agcaggtggg      180
ggcacttctc cctctaacac tctccctgt tgaagctctt tgtgacgggc gagctcaggc      240
cctgatgggt gacttcgcag gcgtagactt tgtgtttctc gtagtctgct ttgctcagcg      300
tcagggtgct gctgaggctg taggtgctgt ccttgctgtc ctgctctgtg acactctcct      360
gggagttacc cgattggagg gcgttatcca ccttccactg tactttggcc tctctgggat      420
agaagttatt cagcangcac acaacanang cagtttccag atttcaactg ctcatca          477

```

<210> 894

<211> 289

<212> DNA

<213> Homo sapien

<400> 894

```

ctgtcttatg gctatgatga gaaatcaacc ggaggaattt ccgtgcctgg ccccatgggt      60
ccctctgggt ctcgtggtct ccctggcccc cctggtgcac ctggtcccca aggcttccaa      120
ggccccctcg gtgagcctgg cgagcctgga gcttcaggct ccatgggtcc ccgaggtccc      180
ccaggtcccc ctggaaagaa tggagatgat ggggaagctg gaaaacctgg tcgtcctggt      240
gagcgtgggc ctctggggcc tcagagtgtc cgaggattgc ccggaacag              289

```

<210> 895

<211> 179

<212> DNA

<213> Homo sapien

<220>

<221> misc_feature

<222> (1) ... (179)

<223> n = A,T,C or G

<400> 895

```

ctggatgggt ccanacaaag tggaatccct ggaaccttta actgagcagt gaaggtcagt      60
gcctcagagc ctgagagatg aacaggacca gagagagagg tgggcaggca ggcacaagg      120
tatgtcttcc tcagactcgg aaccctgtct ttctccacca tccagacgtt cagctacag      179

```

<210> 896
 <211> 557
 <212> DNA
 <213> Homo sapien

<220>
 <221> misc_feature
 <222> (1)...(557)
 <223> n = A,T,C or G

<400> 896
 ccactcactg ctgggaccca ggcacctccc ttctccatcc tctctggatt gtcagtaatg 60
 tcctggaaca gaagcctgtg ggatggcctt gggcacggag aagccctggg gtcagtgtcg 120
 tgcacggatg gcggcagtgt tgaacccagg aggctgaacc cggccccacca cggaagatga 180
 gtgcatggca accgcctgcc ttcacgtcgc tccacttggg aacccccagg tctgggctgt 240
 tctaggtatt gcttcacgtg ccccagcaag cccttaacaa gagggcctgg ttcctgaag 300
 aaccaatccc aggaaggggc cttgatccct ccgccttgct gagagtgaac cctcgtctct 360
 cctcacnctc catttcattt ctgggaattg gggcttagtt tcgaaccttt ggcaaggctg 420
 ttcttactaa tgcccaagcc cctttacccc tctccctata ggttacacag gggagaccag 480
 ggccctcgga gaagactgct gccacacttc cgaatcattc tgcttgccaa ataggtcac 540
 ttcaccagtt gactgac 557

<210> 897
 <211> 495
 <212> DNA
 <213> Homo sapien

<400> 897
 ctggaatctc ctttgcaatc ccatctgata agattaaaaa gttcctcacg gagtcccatg 60
 accgacaggc caaaggaaga gccatcacca agaagaagta tattggatat cgaatgatgt 120
 cactcacgtc cagcaaagcc aaagagctga aggaccggca ccgggacttc ccagacgtga 180
 tctcaggagc gtatataatt gaagtaattc ctgatacccc agcagaagct ggtggtctca 240
 aggaaaacga cgtcataatc agcatcaatg gacagtccgt ggtctccgcc aatgatgtca 300
 ggcaggtcat taaaagggaa agcaccctga acatggtggt ccgcaggggt aatgaagata 360
 tcatgatcac agtgattccc gaagaaattg acccataggc agaggcatga gctggacttc 420
 atgtttccct caaagactct cccgtggatg acggatgagg actctgggct gctggaatag 480
 gacactcaag acttt 495

<210> 898
 <211> 406
 <212> DNA
 <213> Homo sapien

<400> 898
 ccacgactgc atgcccgcgc ccgccagggtg atacctccgc cggtgaccca ggggctctgc 60
 gacacaggga gtctgcatgt ctaagtgcta gacatgctca gctttgtgga tacgcggact 120
 ttgttgctgc ttgcagtaac cttatgccta gcaacatgcc aatctttaca agaggaaacc 180
 gtaagaaagg gccacgcccg agatagagga ccacgtggag aaaggggtcc accaggcccc 240
 ccaggcagag atggtgaaga tgggccaca ggccctcctg gtccacctgg tcctcctggc 300
 ccccttggtc tcggtgggaa ctttgctgct cagtatgacg gaaaaggagt tggacttggc 360
 cccggaccaa tgggcttaat gggacctaga ggccacctg gtgcag 406

<210> 899

<211> 277
 <212> DNA
 <213> Homo sapien

<400> 899
 cctaagagtc attaaaaaat tctccctttg taacctcagt gctggggact gaggcgagcc 60
 ccctcaggtc gctggagtgc accagtcttg gggaagaggt gcaggagaag ctgtgttttt 120
 tatctccaca cgcagtatga agataaaatt acatagtatt acctagacat agacagtatt 180
 acctaggtag atgcaactgc cacctgcacc cttcccagct ctcatttttg ttaggtgatt 240
 tgggataggg atagtgtttt ggggtatggg gggagtg 277

<210> 900
 <211> 389
 <212> DNA
 <213> Homo sapien

<400> 900
 ctgttttgaa atattttactg ttattaaaac ttgcttcaag ggaaattgtg aatatatttc 60
 catatacaag cactagtaac agtaagtggc cctgtcatcc actaactcag gcaaagtaaa 120
 gaatggcatt tttgaaggac attttacctc cccatatgat ttgattggct aggactttct 180
 tctgtaaagt catacctttt cacatcttaa gtttttacat ttgccatttt ccaaattctca 240
 attttgggca agaacgatat agtcacaact atggggctgc tttcaaaagc ggggctccat 300
 ttctactgtc agatcaatgt ggtgctgtaa ccatcttttt atccctacct tcaagaacct 360
 ccttatatga agcctgtctt tatccatca 389

<210> 901
 <211> 453
 <212> DNA
 <213> Homo sapien

<400> 901
 ctggagacac ccacttgggt ggagaagatt ttgacaaccg aatgggtcaac cattttattg 60
 ctgagtttaa gcgcaagcat aagaaggaca tcagtgagaa caagagagct gtaagacgcc 120
 tccgtactgc ttgtgaacgt gctaagcgta ccctctcttc cagcaccag gccagtattg 180
 agatcgattc tctctatgaa ggaatcgact tctatacctc cattaccgt gcccgatttg 240
 aagaactgaa tgctgacctg ttccgtggca ccctggaccc agtagagaaa gcccttcgag 300
 atgccaaact agacaagtca cagattcatg atattgtcct ggttgggtgt tctactcgta 360
 tccccaagat tcagaagctt ctccaagact tcttcaatgg aaaagaactg aataagagca 420
 tcaaccctga tgaagctgtt gcttatgggt cag 453

<210> 902
 <211> 293
 <212> DNA
 <213> Homo sapien

<400> 902
 cctccggccg cccccacggc tcccatggcc tcttctctgc ctaccgtgtg gaggcctaa 60
 ccctgcgtgg catcaatagc ttccgccagt acaagtatga cctgggtggca gtgggcaagg 120
 ctttgagggg catgttccgc aagctcaacc acctcctgga gcgcctgcac cagtccctct 180
 tctctactt gctccccggc ctctcccgct tegtctccat tggcctctac atgcccgtg 240
 tgggtctctt gctcctggtc cttggtctca aggtctgga actgtggatg cag 293

<210> 903
 <211> 228

<212> DNA

<213> Homo sapien

<400> 903

ctggagactc	tgggccagga	gaagctgaag	ctggaggcgg	agcttggcaa	catgcagggg	60
ctggtggagg	acttcaagaa	caagtatgag	gatgagatca	ataagcgtac	agagatggag	120
aacgaatttg	tcctcatcaa	gaaggatgtg	gatgaagctt	acatgaacaa	ggtagagctg	180
gagtctcgcc	tggaaagggt	gaccgacgag	atcaacttcc	tcaggcag		228

<210> 904

<211> 388

<212> DNA

<213> Homo sapien

<400> 904

ccaagcgctc	agatcggcaa	ggggcaccag	tcttgatctg	cccagtgcac	agccccacaa	60
ccaggtcagc	gatgaaggta	tcttcagtct	ccccgaacg	atgaggcacc	atgacgcccc	120
aaccattggc	ctgggccagc	ttgcacgcct	gaagagactc	ggtcacggag	ccaatctggt	180
tgactttgag	caggaggcag	ttgcaggact	tctcgttcac	ggccttggcg	atcctctttg	240
ggttggtcac	tgtgagatca	tccccacta	cctggattcc	tgcactggct	gtgaacttct	300
gccaaagctcc	ccagtcatcc	tggtaaagg	gatcttcgat	agacaccact	gggtagtcct	360
tgatgaagga	cttgtacag	tcagccag				388

<210> 905

<211> 272

<212> DNA

<213> Homo sapien

<220>

<221> misc_feature

<222> (1)...(272)

<223> n = A,T,C or G

<400> 905

ccggagccca	cggnggtcat	ggctgccaga	gcgctctgca	tgctggggct	ggtcctggcc	60
ttgctgtcct	ccagctctgc	tgaggagtac	gtgggcctgt	ctgcaaacca	gtgtgccgtg	120
ccagccaagg	acagggtgga	ctgcggctac	cccatgtca	cccccaagg	gtgcaacaac	180
cggggctgct	gctttgactc	caggatccct	ggagtgcctt	ggtgtttcaa	gcccctgcag	240
gaagcagaat	gcaccttctg	aggcacctcc	ag			272

<210> 906

<211> 525

<212> DNA

<213> Homo sapien

<400> 906

ctgtgcaccc	gagtgtcctt	tccccctaa	gctggcacat	aggagcaaaa	gttcactaac	60
cctgcagtgg	aaggcaccaa	ttgacaacgg	ttcaaaaatc	accaactacc	ttttagagtg	120
ggatgagggg	aaagaaatag	tggtttcaga	cagtgtctct	tcgggagcca	gaagcactgc	180
aagttgacaa	agctttgtcc	ggcaatgggg	tacacattca	ggctggccgc	tcgaaacgac	240
attggtacca	gtggttatag	ccaagagggt	gtgtgctaca	cattaggaaa	tatccctcag	300
atgccttctg	caccaaggct	ggttcgagct	ggcatcacat	gggtcacgtt	gcagtggagt	360
aagccagaag	gctgttcacc	cgaggaagtg	atcacctaca	ccttggaat	tcaggaggat	420
gaaaatgata	accttttcca	cccaaaatac	actggagag	atttaacctg	tactgtgaaa	480

aatctcaaaa gaagcacaca gtataaattc aggctgactg cttct 525

<210> 907

<211> 365

<212> DNA

<213> Homo sapien

<400> 907

gtaaatttta	agtctttcag	ttttatagat	acggaaaaca	agggtgactc	tttaccacag	60
gatgaataaa	gaactaagta	atatgggaaa	tgcagcaatt	tctggactag	ctgagccgat	120
tccttctctgt	gagcacactg	taagctttca	agttctctgg	gcaggaatta	cagcacctgt	180
ccccgtcaat	ggccctgctg	tgtgatgctc	atcgcttccc	tctgtgctgg	agcagtcccc	240
caggtgtcca	tctcctatct	ttttgttcca	atcttctgtg	agttccagct	agcaggcttt	300
acatctgggg	aaaggaaaac	caggggtttt	agctctgttc	tctgctccca	tccttcgctc	360
accag						365

<210> 908

<211> 608

<212> DNA

<213> Homo sapien

<220>

<221> misc_feature

<222> (1)...(608)

<223> n = A,T,C or G

<400> 908

cggaggtgcc	tcagccatgg	catggatccc	tctcttctc	ggcgtccttg	cttactgcac	60
aggacgtgcg	gcctcctttg	aggtgaccca	gccaccttca	atgtccgtgt	ccccaggaca	120
gacagccaag	atcacctgca	ctggagatag	gttgggggat	gaatatgttt	gctgggtatca	180
acagaagcca	ggccagtccc	ctgtattgat	aatataatttg	gataacaagc	ggccctcggg	240
gatccctgac	cgattctctg	cctacgcctc	tgggaacaca	gccactctga	tcatcagcgg	300
ggcccaagtt	atggatgagg	cttattatta	ctgtcaggcg	tgggacggca	gaactgtggt	360
gttcggcgaa	gggaccaacc	tgaccgtcct	aggtcagccc	aaggetgccc	cctcgggtcac	420
tctgttcccg	ccctcctctg	aggagcttca	agccaacaag	gccacactgg	tgtgtctcat	480
aagtgacttc	taccggggag	ccgtgacagt	ggcctggaag	gcagatagca	gccccgtcaa	540
ggcgggagtg	gagaccacca	caccctccaa	acaaagcaac	aacaagtacg	cggnacagcag	600
ctatctga						608

<210> 909

<211> 513

<212> DNA

<213> Homo sapien

<400> 909

ctggtctcaa	actcctcacc	tcaactgatc	cgcccacctt	ggcctcccaa	agtgtctggga	60
ttataggtgt	gagccaccgt	gccc aaagtt	aagtattttt	gatcaagtgt	tttgtctttt	120
gtgcaaggca	tttgtggctc	tgtcatagca	gaggaaaaca	aaacatgcct	atcaaatgaa	180
tcaagtccga	cctcttctca	tattgagcaa	ctagaggctc	aggaacattt	cccctacctg	240
tcattctcat	ctggcatacc	aggtgtacat	actccttctt	attctcctct	gttaccaaga	300
tgttggtccc	attgggtttg	aggtcacgaa	ctccacaaac	tccaaactct	tggacctcag	360
tgtgtgaagg	gaggtcatag	cctagtgtgg	agacatcatt	ttccagcaga	taaaccagac	420
cttggtagaa	gtggtaattc	tcactctcca	tatctgtata	tctgactgac	ttgcccaaga	480
tgtgtttgta	aaaggatcga	gtaaagtagc	act			513

[illegible]

<210> 914
 <211> 252
 <212> DNA
 <213> Homo sapien

<400> 914
 ccaagctggg ggtgcgac tgtggaagaa ctggaggccc ggtgtcatga gcagaggctg 60
 taccctagat gcccgcccca gtgccagcca acccaagaca ggagaaagag tttggcagtt 120
 tcgcctctga ggaatacatg cctggccctc ctgtgaggtg aggcggtagg ggggaaggcg 180
 caggctccga agtctgaggg cttgccggag ggggagtttc tgagcctttt gcatgggtgc 240
 atgccccctg cc 252

<210> 915
 <211> 234
 <212> DNA
 <213> Homo sapien

<400> 915
 ccactgggac tttggcttcc tgatgccgat tgtggatttc tgctgcaaag acagtgatgt 60
 tgagccaggc tgtttcctct ctatccagag gttttgtagt ttttaataaaa ccatcctctg 120
 gattaatagt gaaaaatctg tcgaggtcag tgtgacgata gatggaatac cttatcgggc 180
 tgttggcagc atcagggtct ttggcatgca ctctcccaac cacgggtgcc gcag 234

<210> 916
 <211> 366
 <212> DNA
 <213> Homo sapien

<220>
 <221> misc_feature
 <222> (1)...(366)
 <223> n = A,T,C or G

<400> 916
 ccattcagtc tcanttcaga aaattccaga agaagaaggc tgggtctcag tcctagtggg 60
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 aaagaacaac accctagaga gaagtcaccc acacacaatc cacacacgca tagcaaacct 180
 ccaatgcatg tacagaaacc tgtgatattt atacccttgt aggaagggtat agacaatgga 240
 attgtgagta gcttaatctc tatgtttctc tccattttca ttctctctgc aactattttc 300
 cttgatgttg taataaaatg aagttacgat gagtgatnaa aaaaaaaaaa aaaaaaaaaa 360
 aaaaaa 366

<210> 917
 <211> 492
 <212> DNA
 <213> Homo sapien

<400> 917
 ggcacagcga gggcagcctc tggaggagct ctgcagcctc cacacctacc acgacctccc 60
 agggctgagc tcaggaaaaa ccagccactg ctttacagga caggggggtg aagctgagcc 120
 ccgcctcaca cccaccccca tgcaactcaa gattggattt tacagctact tgcaattcaa 180
 aattcagaag aataaaaaat gggaacatac agaactctaa aagatagaca tcagaaattg 240
 ttaagttaag ctttttcaaa aaatcagcaa ttccccagcg tagtcaaggg tggacactgc 300

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acgctctggc atgatgggat ggcgaccggg caagctttct tcctcgagat gctctgctgc 360
ttgagagcta ttgctttgtt aagatataaa aaggggtttc tttttgtctt tctgtaagg 420
ggtcttccag cttttgattg aaagtcctag ggtgattcta tttctgctgt gatttatctg 480
ctgaaagctc ag 492

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<210> 918

<211> 557

<212> DNA

<213> Homo sapien

<220>

<221> misc_feature

<222> (1)...(557)

<223> n = A,T,C or G

<400> 918

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ctgctcctgg gtaggcgtgc gggccatata gtaggggtag gatactagcc gctcgccgcc 60
gttcagattt gctcccagca cgaagggtt cttctccatc caggcaatga tggcccggac 120
ctccgtggat accgtggcat ctggcgaaag gtagcggtca gggatgggca agttattgtt 180
ggggaccggg taggggacc atttctctc ctcagctccc cagagcacag agttgagatc 240
cgggaaatct tcaaagatgt caaagccctc ctcagtccac agtcccagcg cccagttccc 300
aaactctgag cccatctgcg ctgccacctc gtagccatca gggttcagtg agggcaccag 360
gtggatgcgt gtgtcctgca ccaggctgcg cacacgtggg ttcccatcgc ggtactctcg 420
gcacaggtac tgcattgagc gcagcaacag ctctcgcccc agcacctcgt tgccatggat 480
cccagcagtg tagcggaact cgggctcccc cagttcatgc tcccanggt tgtctgagat 540
ctccatggca tagatct 557

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<210> 919

<211> 407

<212> DNA

<213> Homo sapien

<400> 919

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ccttatgact acaacggccc acgagaaaaa tatggaatcg ttgattacat gatcgagcag 60
tccgggcctc cctccaagga gattctgacc ctgaagcagg tccaggagtt cctgaaggat 120
ggagacgatg tcatcatcat cgggggtctt aagggggaga gtgaccagc ctaccagcaa 180
taccaggatg ccgctaacaa cctgagagaa gattacaaat ttcaccacac tttcatcaca 240
gaaatagcaa agttcttgaa agtctcccag gggcagttgg ttgtaatgca gcctgagaga 300
ttccagtcta agtatgagc ccggagccac atgatggacg tccagggtc caccagggac 360
tcggccatca aggacttcgt gctgaagtac gcctgcccc tggttgg 407

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<210> 920

<211> 340

<212> DNA

<213> Homo sapien

<220>

<221> misc_feature

<222> (1)...(340)

<223> n = A,T,C or G

<400> 920

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cctcttgggc agcnnagggc cctgcctctg tttcatgatg catgggtcat ttgtcttggg 60
tgtcctatcc catatggaga agaaaggggc tctaagttct ggctcttctt tctttggggt 120

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tctctgtacc	tgaggaaacc	aggccctggg	tgactttgca	gatctgctca	ccctcgggtga	180
gcaacagtgt	cagccatgca	agcaggacag	aatgggtgact	gggtgccctt	ggtgagctgt	240
gtatttccta	ggaggtagaa	aactgtggga	aactgtggct	aataaaaaact	aagtgtgagc	300
gtcnaaaaaa	aaaaaaanna	aaaanaaaaa	aagcttgtag			340

<210> 921
 <211> 571
 <212> DNA
 <213> Homo sapien

<400> 921						
ggaaaaataa	ttttattcct	caaatgatca	gcacattcag	aagcaggaca	gaggagctct	60
gatgacatct	ctgggggact	caaagcggcc	ctcattttct	ggtattttcc	caggtgattc	120
tcttccaacc	tgtgagtcct	gctctctttc	ctcccatctg	aagtttgaga	catcctctgc	180
cacaaggaaa	gccaccaata	ccagcccaaa	gagccaccag	agaggaacca	aaccacatgc	240
atcaagttat	aggaaggatg	caagaaggga	aattaggaag	gaaagggagg	agtttagttg	300
gcattctggg	gcatgctaac	atgagggcga	tggctctctc	ccaagtcgct	ggacatatcc	360
cttttctttc	caggtgctcc	aactccaatt	gcagtttgga	ggaacgtgtg	aaacttgttg	420
aagtctcgcg	tgtatgtgcc	cagcatgcaa	gtactcagat	taccgcaccg	cttagatctg	480
gggctgtcca	ggctggagcc	ctctctctct	tgtctctgct	ccagctcact	ggccttcac	540
tgcacatagt	cctgcaccag	tgcagccagc	a			571

<210> 922
 <211> 262
 <212> DNA
 <213> Homo sapien

<220>
 <221> misc_feature
 <222> (1)...(262)
 <223> n = A,T,C or G

<400> 922						
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atccacaaac	cctcgactg	ctgcaggga	agggttgga	aacttctcga	tgtactctgc	120
ctgancagct	tccacattct	catgcccttt	gaagatgatc	tccacagcgc	cctttgctcc	180
catgactgca	atctctgngg	tgggccangc	atanttggtg	tcaccacaaa	ngtgccttaga	240
gctcatgaca	tcntaggcac	ct				262

<210> 923
 <211> 234
 <212> DNA
 <213> Homo sapien

<400> 923						
ccactggggac	tttggcttcc	tgatgccgat	tgtggatttc	tgctgcaaag	acagtgatgt	60
tgagccaggc	tgtttcctct	ctatccagag	gttttgtagt	tttaataaaa	ccatcctctg	120
gattaatagt	gaaaaatctg	tcgaggtcag	tgtgacgatc	gatggaatac	cttatcgggc	180
tgttggcagc	atcaggggtc	ttggcatgca	ctctcccaac	cacggtgcca	gcag	234

<210> 924
 <211> 152
 <212> DNA
 <213> Homo sapien

<400> 924
ccaggattga caggccatcc attcacagcc aggagatgct gggccagttc ctccaagagg 60
tctccgtcat ggcagtgatg aaaacctaac aggggtggccc cctgtgccag ctcaggtgac 120
tggagcccga gggcctgaca ggttcccagc ag 152

<210> 925
<211> 400
<212> DNA
<213> Homo sapien

<400> 925
caatatcatg ccaaggaccc aaacaacctc ttcattggtgc gcttggcaca gggcctgaca 60
catttaggga agggcaccct taccctctgc ccctaccaca gcgaccggca gcttatgagc 120
caggtggccg tggctggact gctcaactgtg cttgtctctt tcctggatgt tcgaaacatt 180
attctaggca aatcacacta tgtattgtat gggctgggtgg ctgccatgca gcccgaatg 240
ctggttacgt ttgatgagga gctgcggcca ttgccagtgt ctgtccgtgt gggccaggca 300
gtggatgtgg tgggccaggc tggcaagccg aagactatca cagggttcca gacgcataca 360
accccagtgt tgttggccca cggggaacgg gcagaattgg 400

<210> 926
<211> 521
<212> DNA
<213> Homo sapien

<400> 926
ccacgtccct attttagaaa tgagaggagt gactgcacac aggaaaaatg ccacttttag 60
caattcaaag tggaaaaact tcttttatat aaaaattatc ccaactccca ccccttggct 120
ctcagtgttg catctcccac agaggtaaag ttgtgccatt ttcccacggc tttaaacaaa 180
gcaaaacaaa accaccaatc ctaataacct ccctccctgc cccgtctcca cgctgtgcgg 240
agagggtctt agcccctcag tcggaattct ccttctcctt catgtgcaag aagacgatgc 300
tgaagatgaa gagccccagc atcatggaga aggcgctggc gtagtagggg taggccgagg 360
ggatgaagcg ctcatactgc gtgtgctgga gtggccgcac ggatacctga gtggaagagt 420
acaggtgtgt gtagcctagc cggttgtaat ccactttaaa ctggaataca ccatacacgt 480
cgggcaactt gaactgaaca ctgtatttgc cacctttctt c 521

<210> 927
<211> 520
<212> DNA
<213> Homo sapien

<400> 927
ccaggctagt ctcgaaactc tgacctcagg tgatctgcct gcctcggcct cccaaagtgc 60
tgggattacc ggcgtgagcc accatgcctg gccttacatt ttttaaaatg agggaacaaa 120
tgaataaatg accaccatgt taggggctgg ctctgaacag aattgtaaag tgggccaagc 180
ttgctctcaa ggtcacctta agcccacggt tgctgtgtcc tgccctctca gggtcatttc 240
ccagcctcca ggcacctgtt cacagaggct gcactctggc tcgcctccac ccctccatcc 300
taaggtgctc cgctgactta gaacaggaca gtcagggaga gaatgtgtct caggagggtg 360
gagtcagatg atcacggcct tcctggcatc tgaggggata cagcttcggg tagcaaagtg 420
tgattttccc tgagccccag gaaagcttgg ccttggctcag aatacattga accctgaggg 480
ccagagagtc cctggggcaa gctctgagag ggaggacctc 520

<210> 928
<211> 492

<212> DNA
<213> Homo sapien

<400> 928
ctgagcttttc agcagataaa tcacagcaga aatagaatca ccttaggact ttcaatcaaa 60
agctggaagt ccaccttaca gaaagacaaa aagaaacccc tttttatatac ttaacaaagc 120
aatagctctc aagcagcaga gcatctcgag gaagaaagct tgcccggctc ccatcccatc 180
atgccagagc gtgcagtgtc cacccttgac tacgctgggg aattgctgat tttttgaaaa 240
agcttaactt aacaatttct gatgtctatc ttttagagtt ctgtatgttc ccatttttta 300
ttcttctgaa ttttgaattg caagtagctg taaaatccaa tctctgagtg catgggggtg 360
gggtgtgaggc ggggctcagc ttcaaccccc tgctcctgtaa agcagtggtt ggtttttccct 420
gagcccagcc ctgggaggtc gtggttaggtg tggaggctgc agagctcctc cagatgctgc 480
cctcgctgtg cc 492

<210> 929
<211> 209
<212> DNA
<213> Homo sapien

<400> 929
ttttttcacc atctaacaaa ggcactttat tgcattacca ttcacaatta acagtcaaga 60
acaaataata ataacaaata aaataacttt taagaggaca aggcattaga aataaaaaag 120
gacactaata acatttgtaa aagcttgtagc tggatgtggt tgccccatt tgtgtgtgtg 180
gttgtgtgtg tgtggtgtg tgttggtg 209

<210> 930
<211> 617
<212> DNA
<213> Homo sapien

<400> 930
cgcgctccttt aacaagcccc gttctcaaaa ggctgggggt atttatataa gaacttattc 60
caaagtgact ctaagatcca tggtcccaag atctagtacg ggctattcat gggtctgagg 120
catgtccagc atgcaggcaa acttatctgt tcaaattgag gtaaaacaga caaaaaacac 180
ttaatatataa cagaagctac ataattaaaa ctaaccttct gctgcttatt taagctaatt 240
atgtatttctt accaaacaga gacctcaag tcaatcattt cttttgattt tagttaccac 300
ccccaaatta agcctcttct ttcaaagcca ttattagtta aaaaaaagtt ttaaaatgaa 360
gaaaaatatt ttttccagaa cttgtatttt gtaattagtg tgatgcaatt tctttttatt 420
tttcaaactt agaaataact catgtatggt actatttggt atttttttca gataccaagg 480
aataaccgaca ggattcataa ataggatttt ctgacactgg caggaaagtc tgctaacggt 540
tacaaaatac caaagactct tctttcaagc ttcaaagatg gctgagaatt aacagttatg 600
attagttttt cagtaca 617

<210> 931
<211> 521
<212> DNA
<213> Homo sapien

<400> 931
ccaacaaaat tgggtgaacac atggaagaac atggcatcaa gtttataaga cagttcgtac 60
caattaaagt tgaacaaatt gaagcagga caccaggccg actcagagta gtagctcagt 120
ccaccaatag tgaggaaatc attgaaggag aatataatac ggtgatgctg gcaataggaa 180
gagatgcttg cacaagaaaa attggcttag aaaccgtagg ggtgaagata aatgaaaaga 240
ctggaaaaat acctgtcaca gatgaagaac agaccaatgt gccttacatc tatgccattg 300

gcgatatatt	ggaggataag	gtggagctca	ccccagttgc	aatccaggca	ggaagattgc	360
tggctcagag	gctctatgca	ggttccactg	tcaagtgtga	ctatgaaaat	gttccaacca	420
ctgtattttac	tccttttgaa	tatgggtgctt	gtggcctttc	tgaggagaaa	gctgtggaga	480
agtttgggga	agaaaatatt	gaggtttacc	atagttactt	t		521

<210> 932

<211> 197

<212> DNA

<213> Homo sapien

<400> 932

ccttgtgacc	aattacatat	gattaaaaatt	acttcccaca	ttcacatcca	cagtactcgt	60
ccaccattta	acatctcaac	caaaacgtta	cacatgtgaa	acaatcacta	acaggcaaaa	120
atactaaacc	tgtatatttg	gtattgcaaa	tacacttatg	catgagcaag	caagggattc	180
acagtgagaa	tctacag					197

<210> 933

<211> 610

<212> DNA

<213> Homo sapien

<400> 933

cctcatttta	acaatatctt	ttttttgtct	ttctgcttcc	aaacottatt	tgccaatgta	60
atgcctttat	ataaagttct	tatgatgaat	gaaaaacttt	caagtgtctg	tgccctatta	120
aatgcattat	ttattaattt	aacttctagt	actctcgata	aagagccagt	gaaatgagtt	180
attgagttcc	agggaaaaaa	atgagaacat	aattttgaat	ttattatctc	tctatacaca	240
cacagttcat	aattggatta	catataataa	taatatcaac	aagtctatca	gtatcgaagt	300
tggatactgg	taatttctca	tgtgaggctc	ttgtgtcaca	gtcagcatag	atctctggag	360
catttgtctg	ttgatctttt	ggtggcctca	aacctcatta	agtgggtgtg	gagatgctgt	420
ttctgccatg	tgagaatgtg	atggcagaat	taacacaacc	ccaccagggg	tacaacagag	480
cactttacat	ccaaaggcag	agagggacac	agcaatgcag	aattccagca	cacttaagag	540
gagcaccatg	ccatccagac	ccattaagat	ggacatagtc	ccatgacaat	tatttgagtt	600
gccatagtag						610

<210> 934

<211> 384

<212> DNA

<213> Homo sapien

<400> 934

ctgctaccag	gggagcgaga	gctgactatc	ccagcctcgg	ctaattgtatt	ctacgccatg	60
gatggagctt	cacacgattt	cctcctgcgg	cagcggcgaa	ggtcctctac	tgctacacct	120
ggcgtaacca	gtggcccgtc	tgccctcagga	actcctctga	gtgagggagg	agggggctcc	180
tttcccagga	tcaaggccac	agggaggaag	attgcacggg	cactgttctg	aggaggaagc	240
cccgttggct	tacagaagtc	atgggtgttc	taccagatgt	gggtagccat	cctgaatggt	300
ggcaattata	tcacattgag	acagaaattc	agaaagggag	ccagccaccc	tggggcagtg	360
aagtgccact	ggtttaccag	gcag				384

<210> 935

<211> 125

<212> DNA

<213> Homo sapien

<220>

<220>

<221> misc_feature
 <222> (1)...(192)
 <223> n = A,T,C or G

<400> 938
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 ctntgcctcc attcacagga aaaaggagct gggagcccca tcctaagggt cccagcatca 120
 gccactgga gggcctggaa cagtccanca ctntgtggga aaggagtggg gaggggaatg 180
 ttttaaaaaa aa 192

<210> 939
 <211> 337
 <212> DNA
 <213> Homo sapien

<400> 939
 ccaaaatatt ggaacacaca gaaccaaacc aggtgtgttc tacacctgca tgagtgaagg 60
 atttccacgt agacacctag gaagagcccg catgccctag actcactcca gaggaaggat 120
 tgatttgcaa ccagaaaggg agctgaaaac caccgagctc catggctctt cattcaaaag 180
 ggaaaataat gattccacgt tgcttttttag agttcaaadc aacatctttc tggataaatc 240
 tattttttta caatcttttt attatttgta aaagatatata aaacaactcc catcagtagc 300
 aatacaaggt tatacatttt aaccagattt tctcagg 337

<210> 940
 <211> 362
 <212> DNA
 <213> Homo sapien

<400> 940
 cctgtccaaa cgtgcgcacc aggaccgagg ggagctccct cccaacacct gctaggaatt 60
 gccactttt aaatggatgg ggttttttat ggggtgaacc tctgttaata cttttgtaca 120
 ctctcactac agtttatatt tttataggct attttctcaa ggtgtttcta gattccacat 180
 atctatttta tataacaagt tattatgtta tgtgtgtgac tcccttgtgt gtatctgtgc 240
 cagcctcagc ctccgagttg cttttccctc tggccctgac tctcactgac tcaccgatgt 300
 ggtgtgcagg cccacttctt accccagata gctcggggcg ctgcctgtag tcatgccgac 360
 ag 362

<210> 941
 <211> 216
 <212> DNA
 <213> Homo sapien

<400> 941
 ctggacatct ttccagcccg ggatacctac catcctatga gcgagtaccc cacctaccac 60
 acccatgggc gctatgtgcc ccctagcagt accgatcgta gccctatga gaaggtttct 120
 gcaggtaatg gtggcagcag cctctcttac acaaaccag cagtggcagc cacttctgcc 180
 aacttgtagg ggcattgtgc ccgctgagct gattgg 216

<210> 942
 <211> 324
 <212> DNA
 <213> Homo sapien

<400> 942

ctgattggct	tcaggccccc	tacctctata	aactctacca	gcattactac	ttcctggaag	60
gtcaaattgc	catectatat	gtctgtggcc	ttgcctctac	agtcctcttt	ggcctagtgg	120
cctcctccct	tgtggattgg	ctgggtcgca	agaattcttg	tgtcctcttc	cccctgactt	180
actcactatg	ctacttaacc	aaactctctc	aagactactt	tgtgctgcta	gtggggcgag	240
cacttggtgg	gctgtccaca	gcctgtctct	tctcagcctt	cgaggccagg	gagcctcaaa	300
tcttcagtct	ctcagagacc	acag				324

<210> 943

<211> 597

<212> DNA

<213> Homo sapien

<400> 943

ctgacaaaat	tcttgggtta	ctaggtgtct	ttcagaagct	gattgcatcc	aaagcaaagt	60
accaccaagg	tttttatctt	ctaaacagta	taatagagca	catgcctcct	gaatcagttg	120
accaatatag	gaaacaaatc	ttcattctgc	tattccagag	acttcagaat	tccaaaacaa	180
ccaagtttat	caagagtttt	ttagtcttta	ttaatttgta	ttgcataaaa	tatggggcac	240
tagcactaca	agaaatat	gatggtatac	aacaaaaaat	gtttggaatg	gttttggaaa	300
aaattattat	tctgaaaatt	cagaaggtat	ctggaaatgt	agagaaaaag	atctgtgcgg	360
ttggcataac	caaattacta	acagaatgtc	ccccaatgat	ggacactgag	tataccaaac	420
tgtggactcc	attattacag	tctttgattg	gtctttttga	gttaccgcga	gatgatacca	480
ttcctgatga	ggaacatttt	attgacatag	aagatacacc	aggatatcag	actgccttct	540
cacagttggc	atttgctggg	aaaaaaaagag	catgatcctg	taggtcaa	ggtgaat	597

<210> 944

<211> 359

<212> DNA

<213> Homo sapien

<400> 944

ctggaagagg	aaaaggagat	actgcagaaa	gaactctctc	aacttcaagc	tgcacaggag	60
aagcagaaaa	caggtagtgt	tatggatacc	aaggctcgatg	aattaacaac	tgagatcaaa	120
gaactgaaag	aaactcttga	agaaaaaacc	aaggaggcgag	atgaataactt	ggataagtac	180
tgttccttgc	ttataagcca	tgaaaagtta	gagaaaagcta	aagagatggt	agagacacaa	240
gtggcccac	tgtgttcaca	gcaatctaaa	caagattccc	gagggctctcc	tttgctaggt	300
ccagttgttc	caggaccatc	tccaatccct	tctgttactg	aaaagaggtt	atcatctgg	359

<210> 945

<211> 367

<212> DNA

<213> Homo sapien

<400> 945

caggatctga	agtttggggg	cgagcaggat	gttgatatgg	tgtttgcgtc	attcatccgc	60
aaggcatctg	atgtccatga	agtttaggaag	gtcctggggag	agaagggaaa	gaacatcaag	120
attatcagca	aaatcgggaa	tcatgagggg	gttcggaggt	ttgatgaaat	cctggaggcc	180
agtgatggga	tcatggtggc	tcgtggtgat	ctaggcattg	agattcctgc	agagaagggtc	240
ttccttgctc	agaagatgat	gattggacgg	tgcaaccgag	ctgggaagcc	tgatcatctgt	300
gctactcaga	tgctggagag	catgatcaag	aagccccgcc	ccactcgggc	tgaaggcagt	360
gatgtgg						367

<210> 946

<211> 335

<212> DNA

<213> Homo sapien

<400> 946

ccacagaggt	ggtattacaa	aatatacaaa	gtgggtttctt	tctttacatt	tcatagaaga	60
agcctgcctc	atttccaaat	gagagcacta	gaagcacaaa	tcatgcagac	catttactat	120
ataacttatg	aaaaatgctg	tacagggctg	tgactataga	tatagagtat	ttggctctgt	180
ttgggaattg	atatctacaa	gggggagggg	caggggagga	ctgtccgata	tcctgacttg	240
ctgggatggg	ggagaagctg	ggatggggga	ggccccaatc	ttgctgcacg	gctacacca	300
ctcctccttt	cctagacaag	gctggagcgc	actgg			335

<210> 947

<211> 384

<212> DNA

<213> Homo sapien

<400> 947

cctcttgagg	cacatccttt	actgcattgt	ggacagcgag	tgtaagtcaa	gggatgtgct	60
ccagagttac	tttgacctcc	tgggggagct	gatgaagttc	aacgttgatg	cattcaagag	120
attcaataaa	tatatcaaca	ccgatgcaaa	gttccaggta	ttcctgaagc	agatcaacag	180
ctccctgggt	gactccaaca	tgctggtgcg	ctgtgtcact	ctgtccctgg	accgatttga	240
aaaccaggtg	gatatgaaag	ttgccgaggt	actgtctgaa	tgccgcctgc	tcgcctacat	300
atcccaggtg	cccacgcaga	tgtccttcct	cttccgcctc	atcaacatca	tccacgtgca	360
gacgctgacc	caggagaacg	tcag				384

<210> 948

<211> 173

<212> DNA

<213> Homo sapien

<400> 948

ctgtggaggg	gacactgtct	ttgaggcatc	actggttcca	caaagggtag	gggaaggtct	60
tgaggggacca	ccccatgcc	tcattaatca	accagaagct	tggcctggag	cagcagcggg	120
gattccagta	gctgtgggca	tacaggatgc	tagggcggcc	acaaccagg	cag	173

<210> 949

<211> 211

<212> DNA

<213> Homo sapien

<220>

<221> misc_feature

<222> (1) ... (211)

<223> n = A,T,C or G

<400> 949

ccatccacgt	tgnaaacag	aataaaatgg	aaattcacct	tgatcatctac	ccgacattgg	60
ccttctgtg	ccacggcatc	atgggctgcc	tgtatggcct	cattcttttc	aaagcatttt	120
gctctgtctt	caggggacat	tttctctgtt	tcagaaagaa	actgtttcag	aactgatcca	180
tcctcaaate	ccagtttgtc	ttgattattg	g			211

<210> 950

<211> 382

<212> DNA

<213> Homo sapien

<400> 950
 cctcatcggtg agtcaggacg tgggtgaaagc tgcagtggtc gctgtgctct ctccagaaga 60
 attcatgggtc ctgttggact ctgtgcttcc tgagagtgcc catcggctga agtcaagcat 120
 cgggctgac aatgaaaagg ctgcagataa gctgggatct acccagatcg tgaagatcct 180
 aactcaggac actcccgagt tttttataga ccaaggccat gccaaagggtg cccaactgat 240
 cgtgctggaa gtgtttccct ccagtgaagc cctccgccct ttgttcaccc tgggcatcga 300
 agccagctcg gaagctcagt tttacaccaa aggtgaccaa cttatactca acttgaataa 360
 catcagctct gatcggtacc ag 382

<210> 951
 <211> 473
 <212> DNA
 <213> Homo sapien

<220>
 <221> misc_feature
 <222> (1) ... (473)
 <223> n = A,T,C or G

<400> 951
 cctctctgcc aggcaaagga gggagctgcg gctctttgac attaaaccag agcagcagag 60
 atacagcctt ttcctccctc tccatgaact ctggaaacag tacatcaggg acctgtgcag 120
 tgggctcaag ccagacacgc agccacagat gattcaggcc aagctcttaa aggcagatct 180
 tcacggggct attatttcag tgacaaaatc caaatgcccc tcttatgttg gtattacagg 240
 aatccttcta caggaaacaa agcacatttt caaaattatc accaaagaag accgcctgaa 300
 agttatcccc aagctaaact gcgtgttcac tgtggaaacc gatggcttta tttcctacat 360
 ttacgggagc aaattccagc ttcggtcaag tgaacggtct gcgaagaagt tcaaagcgaa 420
 nggaacgatt gacctgtgaa ttctttgccg tctaangcag ttgtttatga cag 473

<210> 952
 <211> 312
 <212> DNA
 <213> Homo sapien

<400> 952
 ctgatgggtc tcatagtcct ctgggatggt gtcattgcag cggtaacgca gggttgccca 60
 gatgatgttc tcctgggaga agcagaagac ccccaagcgg ccaccccgca tggttgtgtc 120
 caagaccacg ttgctgtcgg ccaccagctc agggccctca tagaatcgca ccctgatgta 180
 gccacttggt ggccggtgct gcaggaacca acgataggac ttcttgtcct tccaaccac 240
 gtttcgctgg tccttcaca gcagccgcac ctgagactct gtgtctcctg tatgccacag 300
 agcgttccgc ag 312

<210> 953
 <211> 397
 <212> DNA
 <213> Homo sapien

<400> 953
 cgcgtccact gccgaccctc ttggtttctg aaaccaacct ttcttcctgc tctcctcttt 60
 aagagcaaac cccaacatgt ataaggtcac agcaagtggg agccaggaaa agctgtggga 120
 cccctcattt gagtacatc catatggcat ggagaaagaa aacctctctg ccagaaggaa 180
 ctgaactctg gaagtccata ggaaggtcac catgatcagc agataggaaa gcattgccaa 240
 gggctgtccc tcaagagctt agttttctta gggagaccag aaagacatca gatcctgact 300

```
<210> 954
<211> 304
<212> DNA
<213> Homo sapien
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<210> 955
<211> 156
<212> DNA
<213> Homo sapien
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```
<210> 956
<211> 543
<212> DNA
<213> Homo sapien
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```
<210> 957
<211> 528
<212> DNA
<213> Homo sapien
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<400>	957						
ctgtgatcaa	gatgtattaa	aagaatatga	aagagcatct	gggttattct	agaagttctg		60
tgatcaaaac	atattaaaaa	aaattaaagc	gcctctgggt	tattctagaa	gttctctgggc		120
tttatacttg	gatattttaca	gaggaaagtg	aacttcaagt	tctgccactc	ttcaaaatgg		180
gtgacaggag	aggacgtgat	aggacagtta	aaaaaaaaatt	gatagtcatt	ctctgatgga		240
gtgaagcaag	ctttgtcaac	catcaacaaa	tatgacttca	ttggtcacaa	gccttcgaga		300

gatccaacaa	gatttgagtt	ttaaatacag	aacatatttc	aaacagaacc	agcagagtgc	360
tgatgtatga	atggaattga	ttgctgaagg	cagagagtat	aaagaatctc	aagaaacttt	420
tagtgccatt	ttcattttaat	aagccattgg	tatagcaacc	taaaaacctt	ggctgtgatg	480
acaccaggat	gtgtttatgg	aattgctgca	ggagaacaca	attggcag		528

<210> 958

<211> 451

<212> DNA

<213> Homo sapien

<400> 958

ctgtctgacc	atggggacct	tctgtctgaa	gaggagctgg	atgaatgaga	ctctgggaat	60
catctacaca	ggaccaaacc	caacaggcgc	cctggcaccg	gggaggcggg	tagttgtact	120
ctgcttgtag	agtccttgag	cccagtttac	agatctggag	agcaggaggc	caggacaagg	180
acaaaggctg	gaggatggag	taggacccag	gggctctgcc	atcctaggca	tcattcaagg	240
tcttttatga	agactttaca	gatgtcctct	gtaagtagca	tcgagagtgg	agttcagctc	300
ctttctctac	ttttttttgg	tctgatggca	catatttatt	gttctgtggt	ctaatacacag	360
tgtttctaaa	tgtaaaaagt	gcatatgttg	gtgtagctag	tcccgcgaca	ttgagtcct	420
ctgcatgaag	acactgggct	cctgcatcca	g			451

<210> 959

<211> 158

<212> DNA

<213> Homo sapien

<400> 959

ccagaccaag	gctgctggac	ctatgggaat	attcggtgtg	ctgtagagga	tgtgactgtc	60
ctgggtggact	acacagtacg	gaagttctgc	atccagcagg	tgggcgacat	gaccaacaga	120
aagccacagc	gcctcatcac	tcagttccac	tttaccag			158

<210> 960

<211> 235

<212> DNA

<213> Homo sapien

<400> 960

ctgagcaggg	aatccggccg	gaggaaggag	cagcttaccg	actgcgggtg	ttcaccacag	60
gccaggccct	aatatgcacc	cactagttta	gctcagactc	ctctctacat	atgaatggca	120
aaggcacttt	tgatatacac	tgtaaaatac	actgtatttt	agaatcggaa	tctattttct	180
aatgttcccc	tcaagggctg	agtggcagga	aggttgagga	tgcaggactt	tgcag	235

<210> 961

<211> 375

<212> DNA

<213> Homo sapien

<400> 961

cctggaaaga	aaagggatat	gtccagcgac	ttggagagag	accatcgccc	tcatgttagc	60
atgccccaga	atgccaacta	aactcctccc	tttcttccct	aatttccctt	cttgcatcct	120
tcctataact	tgatgcatgt	ggtttggttc	ctctctgggtg	gctctttggg	ctgggtattgg	180
tggctttcct	tgtggcagag	gatgtctcaa	acttcagatg	ggaggaaaga	gagcaggact	240
cacaggtttg	aagagaatca	cctgggaaaa	taccagaaaa	tgagggccgc	tttgagtccc	300
ccagagatgt	catcagagct	cctctgtcct	gcttctgaat	gtgctgatca	tttgaggaat	360
aaaattattt	ttccc					375

<210> 962
 <211> 409
 <212> DNA
 <213> Homo sapien

<220>
 <221> misc_feature
 <222> (1)...(409)
 <223> n = A,T,C or G

<400> 962
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 aagctggggc ctnggctcct nctcatcaaa tacagatcac tgngaccctg tctcctcca 120
 tgggtgctgt ctctcgggcc ccaactgcccc tgetttctgt ttcttctctc acctcctcct 180
 cccccagctc catgtccagc tcgttgccctg cctctgaggg tgtgtagggt gagccactga 240
 tggaacggca gctaaagaag acgattcgct tgagccgctt gttgtagaag aagtagttga 300
 aggaccagag gctaccatcc tccccgaagg gatctgagtc caagtctggg ttatagctgt 360
 agatgtcaca ttcagccagg cagatctcct cgtccaccgc gttccacag 409

<210> 963
 <211> 163
 <212> DNA
 <213> Homo sapien

<400> 963
 gccatggcgt cctattttcga tgaacacgac tgcgagccgt cggaccctga gcaggagacg 60
 cgaaccaaca tgctgctgga gctcgcaagg tcaacttttca ataggatgga ctttgaagac 120
 ttgggggttg tagtagattg ggaccaccac ctgcctccac cag 163

<210> 964
 <211> 344
 <212> DNA
 <213> Homo sapien

<400> 964
 ccaactggctg agttattggc ctggcaggta tagagtccgc tgttcttctc agtgatgttg 60
 gagataaaga gctcttggtg gtgttgctgg atgttcccat caatcagcca agaatactgt 120
 gcagggtgggt tagaggctgc atggcaggag aggctgaggt tcaccctctg acggtaatag 180
 gtgtatgagg gggaaatggt ggggtcgtct gggccataga ggacattcag gatgactggg 240
 tcgctgtggt caacacttaa ttcgttctgg attccacact catagggtcc tacatcattc 300
 cttgtgacac tgagtagagt gagggtcctg ttgtcattgg acag 344

<210> 965
 <211> 461
 <212> DNA
 <213> Homo sapien

<400> 965
 ctgagctttc agcagataaa tcacagcaga aatagaatca ccctaggact ttcaatcaaa 60
 agctggaagt ccaccttaca gaaagacaaa aagaaacccc tttttatata ttaacaaagc 120
 aatagctctc aagcagcaga gcattctcag gaaggaagct tgcccggctc ccatcccatc 180
 atgccagagc gtgcagtgtc cacccttgac tacgctgggg aattgctgat tttttgaaaa 240
 agcttaactt aacaatttct gatgtctatc ttttagagtt ctgtatgttc ccatttttta 300

ttctttctgaa ttttgaattg caagtagctg taaaatccaa tctttgagtg catgggggtg 360
 ggtgtgaggc ggggctcagc ttcaaccccc tgtcctgtaa agcagtggct ggtttttcct 420
 gagcccagcc ctgggagggtc gtggtagggtg tggaggctgc a 461

<210> 966
 <211> 246
 <212> DNA
 <213> Homo sapien

<400> 966
 cctttcacag aactaccat tgagtgggtt gatgcagggt gcagccttca gtccccgagt 60
 actgggttct gataaaattc cacagaatcc agcatcactg ggctcagacg gcatccactg 120
 tagtaacta tttgtaaatg gggacatata tcccagcac cagtaggaca cattgatctt 180
 ccgaaggccg acccatgggg ttaagggtgag cttggacatg ctctgagatg actgcattat 240
 tcgcag 246

<210> 967
 <211> 244
 <212> DNA
 <213> Homo sapien

<400> 967
 ctggagcatt ggcagggaca gtcagaaagg agacaagtga aaacgggtcag atggacacag 60
 gcggaggaga aaagacagag ggagagagac catcggaac aatcagaggg gccgagacga 120
 tcagaaaagg gtcagcccga gacaggctga gccagagttt ctagaagcag tttccaattc 180
 aacggctcgc tttgagggcc aacgtgtcct aggccgaggc tgcagaagcg ctcacacact 240
 cagc 244

<210> 968
 <211> 436
 <212> DNA
 <213> Homo sapien

<400> 968
 ccaaagtctt taccctatctt aacccttctg atattttctga ctgctcactg ttcataattat 60
 aggggaccag atttgaata tagaattctc cataacatga atgaaattaa tgctgtccaa 120
 gccagcatgg tggcttcata ttaagtagta acagaagtct gaacaattgg ataaatttga 180
 cttccaagac agctaaactt ttcaactgca attttaaaaa ctacactaca ctgttatagt 240
 taatctgaca aaaatgtcct caaagagtac tttattttat ttaaagcatc tgtttaattc 300
 aacctttaat aattttgcaa agaagggtac gtgtgtatct taatatagcc tgacctgaat 360
 ttatatgttt ttagcttttag tatttaactt tttgtaacaa ataaaccttt tttaaaacaa 420
 gtttaaaaaa gaaaaa 436

<210> 969
 <211> 383
 <212> DNA
 <213> Homo sapien

<400> 969
 ctggctccct tgtctccagg gctttggagg atcagggtag ggagggctct gtctctaagc 60
 cagggtgtcag gatcagaatc atgggtagaa ggtgccattc agctcacagc cgcacccaga 120
 atcctttgca gccctccttc tttatttttt tccattgca ttctgggagt ccacatctgg 180
 ctttctcagc cactgttcat caccaggggt tttaggagga aggcttggtc cctgtcttcc 240
 cagacccacc atgcctggag aggtcaggat ggaactacct cattcggcga attagcccca 300

aattgaacgc tgaatcgtgt cccatgagat caggcgccat ctgtaaagtc tcctctggaa 360
atgccaatcc atccttcccc cag 383

<210> 970
<211> 543
<212> DNA
<213> Homo sapien

<400> 970
ctgtagcttt tgtgggactt ccactgctca ggcgtcagge tcaggtagct gctggccgcg 60
tacttgttgt tgctttgttt ggaggggtgt gtggtctcca ctcccgctt gacggggctg 120
ctatctgcct tccaggccac tgtcacggct cccgggtaga agtcacttat gagacacacc 180
agtgtggcct tgttggcttg aagctcctca gaggagggcg ggaacagagt gaccgagggg 240
gcagccttgg gctgacctag gacggtcagc ctggctccctc cgcggaacac cgaagtgcta 300
ctgtttgtat atgagctgca gtaataatca gcctcgtcct cagcctggag cccagagatg 360
gtcagggagg ccgtgttgcc agacttgagg ccagagaagc gattagaaac ccctgagggc 420
cgatcagtga catcataaat catgagtttg ggggctttgc ctgggtgctg ttggtaccag 480
gagacatagt tataaaaacc aacgtcactg ctgggtccag tgcaggagat ggtgatcgac 540
tgt 543

<210> 971
<211> 416
<212> DNA
<213> Homo sapien

<400> 971
ccagactgac ttcaaaaaat taatgtgtat ccagggacat tttaaaaacc tgtacacagt 60
gtttatttgt gttaggaagc aatttcccaa tgtacctata agaaatgtgc atcaagccag 120
cctgaccaac atggtgaaac cccatctgta ctaaacataa aaaaattagc ctggcatggt 180
gggtgtacgcc tgtaatccca gtgacttggg aggctgaggc aggagaatcg cttgaaccgc 240
ggaggcggag gttgcagtga gctaagatcg caccactgta ctccagcctg ggcaacagcg 300
agactccatc tcaaaaaaaaa aggaaatgtg tatcaagaac atgattatcc aggggtattt 360
totaattcag atcatcaaac tgattatata gaagagtgg ctttaaaatg tttgca 416

<210> 972
<211> 242
<212> DNA
<213> Homo sapien

<400> 972
ccaaaaatcc caaaacatca ttttcaatca gtagagaagt gcttaggggtt gaaaattgat 60
ttcatttgct actgaatttg gtaaatcctg ggtaactttt atcaagatga agacatttta 120
ccctacctac tctagaaata tacaacaatg ttatatatta cactccttgg aaacatttga 180
ggaaaaaaat gcaatttgca cttcactttg ttggaatatc ccatagcact caataaactc 240
ag 242

<210> 973
<211> 347
<212> DNA
<213> Homo sapien

<400> 973
cctgcagggg atggaacctt ccagaagtgg ggcgctgtgg tggcgcttc tggagaggag 60
cagagatata cctgccatgt gcagcatgag ggtctgcca agccctcac cctgagatgg 120

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gagctgtctt cccagccac catcccatc gtgggcatca ttgctggcct ggttctcctt 180
ggagctgtga tcaactggagc tgtggctcgt gccgtgatgt ggaggaggaa gagctcagga 240
cattttcttc ccacagatag aaaaggaggg agttacactc aggctgcaag cagtgcagct 300
gccagggct ctgatgtgtc tctcacagct tgtaaagtgt gagacag 347

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<210> 974
<211> 571
<212> DNA
<213> Homo sapien

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<400> 974
gaaagagcga gatgcgagaa cacttttggc taaaaatctc ccttacaaag tcaactcagga 60
tgaattgaaa gaagtgtttg aagatgctgc ggagatcaga ttagtcagca aggatgggaa 120
aagtaaaggg attgcttata ttgaatttaa gacagaagct gatgcagaga aaacctttga 180
agaaaagcag ggaacagaga tcgatgggcg atctatttcc ctgtactata ctggagagaa 240
aggtcaaaat caagactata gaggtggaaa gaatagcact tggagtgggtg aatcaaaaac 300
tctggtttta agcaacctct cctacagtgc aacagaagaa actcttcagg aagtatttga 360
gaaagcaact tttatcaaaag taccacagaa ccaaaatggc aaatctaaag ggtatgcatt 420
tatagagttt gcttcattcg aagacgctaa agaagcttta aattcctgta ataaaagga 480
aattgagggc agagcaatca ggctggagtt gcaaggaccc aggggatcac ctaatgccag 540
aagccagcca tccaaaactc tgtttgtcaa a 571

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<210> 975
<211> 221
<212> DNA
<213> Homo sapien

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<220>
<221> misc_feature
<222> (1)...(221)
<223> n = A,T,C or G

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<400> 975
ctggaggtgc ctcanaaggt gcattctgct tcctgcaggg gcttgaaaca ccaaggcact 60
ccaggggatcc tggagtcaaa gcagcagccc cggttggtgc actccttggg ggtgacatgg 120
gggtagccgc agtccaccct gtcttggct ggcacggcac actggtttgc agacaggccc 180
acgtactcct cagcagagct ggaggacagc aaggccagga c 221

```

```

<210> 976
<211> 316
<212> DNA
<213> Homo sapien

```

```

<400> 976
ccatcagatt gtcacagact tttataaccc tttgatccct accaacgtta agtatgagtt 60
tggccctgcc atcttcattg gctgggcagg gtctgcccta gtcaccttg gaggtgcact 120
gctctcctgt tcctgtcctg ggaatgagag caaggctggg taccgtgcac cccgtctta 180
ccctaagtc aactcttcca aggagtatgt gtgacctggg atctccttgc cccagcctga 240
caggctatgg gagtgtctag atgcctgaaa gggcctgggg ctgagctcag cctgtgggca 300
gggtgccgga caaagg 316

```

```

<210> 977
<211> 335
<212> DNA

```

<213> Homo sapien

<400> 977

cctgtttgtc	tgtacagcaa	tgcagatgcg	caggcccatc	ctggtggagg	acccagatgc	60
aggagcaaa	tattcgggtt	gtgttgctaa	gagtcgcagg	aactactgct	agtgatacta	120
ggcttgctgc	aggaggatgt	cacgctgaga	aaggagatg	actaggagca	gaaaaagtac	180
tctcactgtt	ccagcttcca	gccccatcct	agcagaatga	atgcatttta	aaatcagtcc	240
acattcacat	gtgctgagaa	ggttgttagt	ggcccccat	ctgggcaaag	cagacccaag	300
atggtgctaa	gtgcagagt	cagagcattc	ttgtg			335

<210> 978

<211> 280

<212> DNA

<213> Homo sapien

<400> 978

cctaacaccc	aagctcttcc	ttgcagaaga	gctgagatgc	taaggagacc	atctggagt	60
tcataataag	cccttgggat	ttgctgagct	cccacatggc	tttcttcaac	cacctggccc	120
actttcttca	accacattcc	actttggaat	gcgtgtcttt	aaggcaccaa	gtgatcttaa	180
gaatgggctc	tgtttttgaa	ttcagcaatc	caagttccta	tctatctcgg	tgggacctcc	240
aaaaaaaaa	aaaaggattg	gcttggcttc	taatgtaagg			280

<210> 979

<211> 318

<212> DNA

<213> Homo sapien

<400> 979

ctgtccagat	gacagtaaga	ttccactgtc	tgtaatcctc	atggtgccag	gtctcctggg	60
gcatctaggg	caatgatgct	actgcagttt	atgcagttac	acagtcaagt	ctgtgccaaa	120
ggaggtccca	tccggcggcc	aggtttctgt	tcagtctggg	gagcaatgcc	aactggctgc	180
ccccatagcc	tggcatgagc	tgatggccca	gtgcaatccc	aaagcaaaga	agggcagaac	240
tgggccaaga	agctgtggta	atgtgctctc	cctgcctccg	acagcgtcgt	cctctccttt	300
tgcagcccca	cacgcagg					318

<210> 980

<211> 568

<212> DNA

<213> Homo sapien

<400> 980

ccagcactgg	ctccttgatg	gttttcctag	gacattagga	caagccgaag	ccctggacaa	60
aatctgtgaa	gtggatctag	tgatcagttt	gaatattcca	tttgaaacac	ttaaagatcg	120
tctcagccgc	cgttggattc	accctcctag	cggaagggtg	tataacctgg	acttcaatcc	180
acctcatgta	catggtattg	atgacgtcac	tggtgaaccg	ttagtccagc	aggaggatga	240
taaaccogaa	gcagttgtcg	ccaggctaag	acagtacaaa	gacgtggcaa	agccagtcat	300
tgaattatac	aagagccgag	gagtgtctca	ccaattttcc	ggaacggaga	cgaacaaaat	360
ctggccctac	gtttacacac	ttttctcaaa	caagatcaca	cctattcagt	ccaaagaagc	420
atattgaccc	tgcccaatgg	gagaaccagg	aagatgtggg	cattcattca	atagtgtgtg	480
tagtattggg	gctgtgtcca	aattagaagc	taactgaggt	agcttgcagc	atctcttcta	540
gttgaaatgg	tgaactgata	ggaaaaca				568

<210> 981

<211> 550

<213> Homo sapien

ccatccccct	ttagaacgta	tcttaatgtg	aacataaatt	gttcttcatg	atgcttaaaa	60
gcttacatat	aattttcatt	cttagaaaaa	cgccacattt	tggatcctgg	atttttctga	120
atatcatgat	tgaaaaaaac	aaaacaaaaa	atgaacccaa	atcaaagtg	ggttaaactt	180
atatgagaaa	gatttttcaa	ccagatggtc	attcaaaaaa	gttggagctg	taagtgccgg	240
cgactgagga	cacagggtta	attcctcgct	gctggtgga	ggctagagaa	catcttcaaa	300
agagggtagc	aagacgtgct	cctaggggag	gctcagtg	gtctcgtctg	cccaagcatt	360
ttcagtcctg	cttgggtcaat	gacatcgagt	aagtttttgg	catccacagc	cagggcgtga	420
gcagcagtc	gcatttgctt	tttgactct	tgctggaggc	tggtcatgac	atactgctgg	480
gccagtttca	tcttggtgat	gagctcacc	aggtcagagt	tcaatagctt	ctgtgccatc	540
tcaatctctc						550

<211> 524

<213> Homo sapien

ccaaggctcag	aggctgatgc	aacaggccct	cttctcccca	gggccaggct	cctgtccagc	60
ctgggcactg	cccagagtga	tggcattggg	ccggatgctg	ttctgtctct	gcttggacac	120
cttcgcaaaag	atttctttca	ggacagtctc	aaaggctagc	tcaacattgg	tagagtccag	180
ggctgaggtc	tccaggaaga	gcagtcatt	gttttcagcg	aacattcggg	cctcctcagt	240
gggcacttcc	cgggcctggc	tgaggtcact	tttgttacc	acgagcatga	cgacgatcgt	300
ggcttcagca	tggtcataga	gctccttcag	ccatcgctcc	accacagcat	aggtctggtg	360
cttggttagg	tcaaacacca	ggagggcccc	cactgcacca	cgatagtacc	cttgaagaca	420
aagttataat	cttcctcagt	tccattcccc	atcttggtct	cgcattggagg	gtgcagggtgt	480
cttcggggac	agagggcgaca	aatctgtgtg	ttggctcaat	gccc		524

<211> 140

<213> Homo sapien

```

ccttcgtgtgcc ctaacagcca gtccctgttt aaagtggaag agacctgtgg ctgccgttgg      60
acctgccccct gtgtgtgcac aggcagctcc actcggcaca tcgtgacctt tgatgggcag      120
aatttcaagc  tgactggcag                                     140

```

<211> 358

<213> Homo sapien

tggagcggcc	gcccggcagg	tccaacgagt	cacaacagtg	caataggtag	aggattaaaa	60
actgcatcaa	acaggtgctg	aaaataaata	ctacctagga	gaaggagggtg	agagccctcg	120
tgtggggttt	gttttcgacc	ccttgagtg	gtgtgggggt	tgtcttcga	gccacgagcc	180
tggcctgtct	cgcggtgctg	ttcactctga	cagagtgcgc	ctgcagcacg	ttgcctccag	240
ggcccagcct	cccagaagcc	tcagagcatc	agagcatccg	tcccatcgga	tggaccagaa	300
acaagaaaat	gggggtgggt	gaatcacagc	tatcattcaa	aggaaaggaa	tttttttc	358

<210>	989
<211>	193
<212>	DNA

<213> Homo sapien

<400> 989

ccagccgtgt	cccagacttg	tagtttgatc	ttcttccct	ctatatccac	agtgcggatc	60
ttgaaatcaa	ttccgatgg	ggagatgtaa	gtgttggtga	agttgtctc	tgcaaagcga	120
atgatcagac	aagtcttgcc	cacccccgag	tccccgatca	gcagcaactt	gaagaggtgg	180
tcgtaggctt	tgg					193

<210> 990

<211> 499

<212> DNA

<213> Homo sapien

<400> 990

cctcaaccaa	gagggttgat	ggcctccagt	caagaaactg	tggctcatgc	cagcagagct	60
ctctctcct	ccagcaggcg	ccatgcaagg	gcaggctaaa	agacctccag	tgcatcaaca	120
tccatctagc	agagagaaaa	ggggcactga	agcagctatg	tctgccaggg	gctaggggct	180
cccttgacga	cagcaatgct	acaataaagg	acacagaaat	gggggaggtg	ggggagccct	240
atthttataa	caaagtcaaa	cagatctgtg	cgttcattcc	cccagacaca	caagtagaaa	300
aaaaccaatg	ctgtggtttc	tgccaagatg	gaatattcct	cctcctagtt	ccacacatgg	360
cgtttgcaat	gctcgacagc	attgcactgg	gctgctgtct	ctgtgttctg	gcaccagtag	420
cttgggcccc	atatacactt	ctcagttccc	aacaagggtc	tatgggccga	ggggcaggct	480
ccaattttca	agcacacga					499

<210> 991

<211> 262

<212> DNA

<213> Homo sapien

<400> 991

ctgccagcca	ggctgtggtc	agtcctctgg	caggcaatct	tccggaccga	gagcctctgt	60
ccattagtgt	cagccccgag	ggggccacga	cggaggccgc	ccaatgtcca	ctgtgatatt	120
ggtgaagagt	ggttgccgag	acacctccaa	gacctggtac	cgcactgacc	caatgccgtc	180
ccgcttcatg	gtcagcttcg	tgttttgaat	cttggtaaac	ctctgagggg	taggttcgtt	240
atgcttctgc	cggctgtgct	tg				262

<210> 992

<211> 535

<212> DNA

<213> Homo sapien

<220>

<221> misc_feature

<222> (1)...(535)

<223> n = A,T,C or G

<400> 992

ctgctgcttg	tgaaattcat	gtgtggtact	aagtacctta	catgaattat	ttcattttaac	60
cctoccaaaca	gtctcctttg	tacgtgctgn	ncctctctgcc	tggaaacact	gtttcccacc	120
cccaaccccc	aattcttctg	tttatttttc	ttgagacaga	gtctcactgt	gtagcccaga	180
ctggagtgc	gtggcgcat	ctcggctcac	tccaatctcc	gcctcccggg	tccctgttca	240
agcagttctc	ctgcctcagc	ctcctgagta	gctgggatta	caggcacacg	ccaccatgtc	300
cagetaattt	ctgtattttt	agtagagatg	gggtttcacg	atgttggtta	ggatgggtctc	360
gatctctggt	cagagtcctt	tctgtaaata	tccttggtta	agaagcaatt	ttagactgta	420

gctgttgcaa atgctttaag gaagaagcaa aacaactgtc agtcttnctg aaatgaagaa 480
actacaccag ggctgctata tcagagcaac cccaaccagc actncaatca tgatg 535

<210> 993
<211> 232
<212> DNA
<213> Homo sapien

<400> 993
ctgctgctct cccctcccag tctctactca ctgggatgag gttagggtcat gaggacacca 60
aaaacctaata aataaaca aaagccaaaca agccttagct tttcttaaag gctgaaatgc 120
ctggaagtgt ccttttattt ataaaataac ttttgtcata tttcttatac atgtttcttg 180
taagaaattc agaaactaca gacaaagaga gtggaaatta cccactgtca gg 232

<210> 994
<211> 203
<212> DNA
<213> Homo sapien

<400> 994
ccagcagatc atccacgacg accaccctct gtccctggctc cagggcgctct ttctgaatct 60
ccagctcagc cttcccgtac tccaggggaat aggaggccca cagagtgggg cctggcagct 120
tcccccgctt tcggatgagc acgcagccca gtccaagctc ctggggccagg gaggggcca 180
agaggaagcc tcgggagtct agg 203

<210> 995
<211> 238
<212> DNA
<213> Homo sapien

<400> 995
ccatgcctgc cccgcccact ctgtatatat gtaagttaaa cccgggcagg ggctgtggcc 60
gtctttgtac tctgggtgatt tttaaaaatt gaatctttgt acttgcatg attgtataat 120
aattttgaga ccaggtctcg ctgtgttgct caggctggct ccaaactcct gagatcaagc 180
aatccgcccc cctcagcctc ccaaagtgtc gagatcacag gcgtgagcca ccaccagg 238

<210> 996
<211> 379
<212> DNA
<213> Homo sapien

<400> 996
ctgcagcctg ggactgaccg ggaggctctg accatttacc caccacaggt aggttgtgtt 60
ctgaacctca ggttcacagg tgaaggccac agcatccttg tcctccacgg ggttgaggtt 120
gttgctggag atggagggct tgggcagctc cgggtataca tggaactgtc cggttgcttc 180
ttcattcaca agatctgact ttatgacttg tagggatatag aatcctgtgt cattctgggt 240
gacgttctgg atcagcaggg atgcattggg gtatattgtc tctcgaccac tgtatgcggg 300
ccctggggta gcttggttag ttctattac atatcctaca attagactgt tgccatccac 360
tctttcgctt ttgtaccag 379

<210> 997
<211> 210
<212> DNA
<213> Homo sapien

```

<400> 997
ccatccgaag caagattgca gatggcagtg tgaagagaga agacatattc tacacttcaa      60
agcttttggtg caattcccat cgaccagagt tggttccgacc agccttggaagggtcactga      120
aaaatcttca attggattat gttgacctct accttattca ttttccagtg tctgtaaagg      180
ccgtggagaa gtgtaaagat gcaggattgg      210

<210> 998
<211> 207
<212> DNA
<213> Homo sapien

<220>
<221> misc_feature
<222> (1)...(207)
<223> n = A,T,C or G

<400> 998
ggtggctgtg ctggggggcgc cccacaaccc tgctcccccg acgtccaccg tgatccacat      60
ncgcagcgag acctccgtgc cgcaccatgt cgtctgggtcc ctgttcaaca ccctcttcat      120
gaacccttgc tgcctgggct tcatagcatt cgctactcc gtgaagtcta gggacaggaa      180
gatggttggc gacgtgaccg gggccca      207

<210> 999
<211> 315
<212> DNA
<213> Homo sapien

<400> 999
ccaatgggct ttgctgtagc ttgctgaaat caccaagcag gagagattta accagaggcg      60
atgtgtccag tcaccagcat agagccatcc tctgtgtcac catccacacg cagggccttc      120
tggcagacct catgcaatgc cctccatggt aatattcatc agaaaatgga taattagggg      180
ggccagcaaa aatatcaagg gtcaaataac gcacatttct gtttaggccca tctatggctt      240
tcattctctc tgaagtcaac tggaattcaa acacctgcac gttctgtctg atgcgctgct      300
cattgtagct cttgg      315

<210> 1000
<211> 186
<212> DNA
<213> Homo sapien

<400> 1000
ctgttactca agaagatgta tttaatgctt gacaataaga gaaaggaagt agttcacaaa      60
ataatagagt tgctgaatgt cactgaactt acccagaatg ccctgattaa tgatgaacta      120
gtggagtggg agcggagaca gcagagcgcc tgtattgggg ggccgccc aa tgcttgcttg      180
gatcag      186

<210> 1001
<211> 173
<212> DNA
<213> Homo sapien

<400> 1001
ccacaaaagcg gaaactcatc cacttttggc tttttccgcc ccaggtcaaa aatgcgaatc      60

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```
<210> 1002
<211> 302
<212> DNA
<213> Homo sapien
```

```
<210> 1003
<211> 368
<212> DNA
<213> Homo sapien
```

```
<210> 1004
<211> 294
<212> DNA
<213> Homo sapien
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```
<210> 1005
<211> 414
<212> DNA
<213> Homo sapien
```

<400> 1005							
ctgaagcact	cttcagagac	tacgtccaca	gacactgatg	ctgaggcctt	tcttgtaagt		60
gaagaaaaaag	gaatgcagca	aagaagagtt	cgacattgga	gtccttagtt	ccatcaggat		120
cccattcgca	gccttttagca	tcatgtagaa	gcaaactgca	cctatggctg	agataggtgc		180
aatgacctac	aagattttgt	gttttctagc	tgtccaggaa	aagccatctt	cagtcttgct		240
gacagtcaaa	gagcaagtga	aaccatttcc	agcctaaact	acataaaaagc	agccgaacca		300
atgattaaaq	acctctaagg	ctccataatc	atcattaaat	atgccccaaac	tcattgtgac		360

ttttttatttt atatacagga ttaaaatcaa cattaaatca tcttatttac atgg 414

<210> 1006

<211> 272

<212> DNA

<213> Homo sapien

<400> 1006

cgggagccca	cggtgggtcat	ggctgccaga	gcgctctgca	tgctggggct	ggtcctggcc	60
ttgtgtcct	ccagctctgc	tgaggagtac	gtgggcctgt	ctgcaaacca	gtgtgccgtg	120
ccagccaagg	acaggggtga	ctgcggctac	ccccatgtca	cccccaagga	gtgcaacaac	180
cggggctgct	gctttgactc	caggatccct	ggagtgcctt	gggtgtttcaa	gcccctgcag	240
gaagcagaat	gcaccttctg	aggcacctcc	ag			272

<210> 1007

<211> 313

<212> DNA

<213> Homo sapien

<220>

<221> misc_feature

<222> (1) ... (313)

<223> n = A,T,C or G

<400> 1007

cctgccttac	tctnttcct	ttccccaggg	actcttggtt	ttcagaagcc	cctctggaat	60
gtectacctg	gcctaacccc	ataccagcag	tgcagacaag	gaggcactcc	tactatagtg	120
gggtccagccc	atggagagac	tcacttcctg	ccccaacacc	tcttccccta	gacctgagg	180
gccaggacaa	tgtcttagtg	ccttccaact	tggcagagtg	aggcccatg	agacagagag	240
aaagggggaa	gagggaaata	cctttatcca	aataaatacc	catccaaaat	tattttgtgat	300
aggtgaaaaa	tgg					313

<210> 1008

<211> 317

<212> DNA

<213> Homo sapien

<400> 1008

cctcaatgtc	gtgctagagg	ggccgaagaa	ggccgtgaac	gacgtgaatg	gcctgaagca	60
atgtttggca	gaattcaagc	gggatctgga	atgggttgaa	aggctcgatg	tgacactggg	120
tccggtaccg	gagatcggtg	gatctgaggc	gccagcacct	cagaacaagg	accagaaagc	180
tgttgatcca	gaagacgact	tccagcgaga	gatgagtttc	tatcgccaag	cccaggccgc	240
agtgtttgca	gtcttaccct	gcctccatca	gctcaaagtc	cctaccaagc	gacccactga	300
ttattttgcg	gaaatgg					317

<210> 1009

<211> 456

<212> DNA

<213> Homo sapien

<400> 1009

tttttttgta	gggtatagaa	aatacathtt	taattttgat	agagttcaca	aatgacagca	60
ttgacatttc	tttaaacaaa	tacttctgtc	aaggcacagc	attaccatgt	gtccccagat	120
gccaagagg	cagtgatctc	atgtccccct	gaggttttagc	agagccacca	atgtcaatag	180

```

ggtggctgac ggggcctaga tttgctacca gataagccaa tgagacatgc tgtcagattt 240
atggttacat aatcaagtat ttaaaaagat gcacaatagg taactgcaat gagcttgctc 300
tgcatttagc gatagttcct ttcaaacaaa gaagatagtt ttcagtatca agaaggatgc 360
ctatatgtat gtcttccatg gagcctttcc tacaaattgc tttcattaca cattaacagg 420
agttcagctt tattgtgacc ttcttgagtc attcag 456

```

<210> 1010

<211> 196

<212> DNA

<213> Homo sapien

<400> 1010

```

ctgggcatgg gctgaggaga ggtcttgctt gcccccttca actttccatc tcagaactat 60
aaactgctag gctgcaagga gagaagggct aagtgggggt cagacaggag agaagggcag 120
gaggcagtga gccccgatga cccaccaact ccaccaggcc ctgacaggga agcccccttg 180
gtagtatca ttttgg 196

```

<210> 1011

<211> 449

<212> DNA

<213> Homo sapien

<400> 1011

```

ccttgcggt gctgcgaaag gccacggcgc tgcctgcccg ccgggcccag taactttgatg 60
gttcagagcc cgtgcagaac cgcgtgtaca agtcaactgaa ggtctgggtc atgctcgccg 120
acctgaagga gagcctcggc accttccagt ccaccaaggc cgtgtacgac cgcctcctgg 180
acctgcgtat cgcaacaccc cagatcgtca tcaactatgc catgttcttg gaggagcaca 240
agtacttcga ggagagcttc aaggcgtacg agcgcggcat ctgctgttc aagtggccca 300
acgtgtccga catctggagc acctacctga ccaaattcat tgcccgtat gggggccgca 360
agctggagcg ggcacgggac ctgtttgaac agyctctgga cggctgcccc ccaaaatatg 420
ccaagacctt gtacctgctg tatgcacag 449

```

<210> 1012

<211> 289

<212> DNA

<213> Homo sapien

<220>

<221> misc_feature

<222> (1) ... (289)

<223> n = A,T,C or G

<400> 1012

```

ccaggaccac aacccccacgc ttagctggt agcgcagggc aatcagggct ggggttcgct 60
tgtgttttt tgccaaggca caaaggactg ggtcctccaa gagcaccggg gagttcgggt 120
ccacccatgg ttcttctcgg tgggatccca gagcactata ggcaaccaga acaatgtctt 180
ttgacttgca gaaatccagc agttttctct ggttgaagta aggatgacat tccacctggt 240
tgcagacagg ctgtgacttg agccctggct tgnnaggat catctccag 289

```

<210> 1013

<211> 221

<212> DNA

<213> Homo sapien

```

<220>
<221> misc_feature
<222> (1)...(221)
<223> n = A,T,C or G

<400> 1013
tctgtaaatg ctgcgttcct aatttagtaa aataaaagaa tagacactaa aatcatgttg 60
atctataatt acacctatgg gatcaataag catgtcanna ctgattaatg tctactgtaa 120
aaatttggtg gnnaaatttt catttgatat tagatataaa tatctgaata taaataattn 180
taatatacta gtcgatgatg gtgttgatt ttaaaaatta t 221

<210> 1014
<211> 512
<212> DNA
<213> Homo sapien

<400> 1014
gggccccga agcctctaca atgggctggt tgccggcctg cagcgccaaa tgagctttgc 60
ctctgtccgc atcggcctgt atgattctgt caaacagttc tacaccaagg gctctgagca 120
tgccagcatt gggagccgcc tcctagcagg cagcaccaca ggtgccctgg ctgtggctgt 180
ggcccagccc acggatgtgg taaaggctcg attccaagct caggccccgg ctggagggtg 240
tcggagatac caaagcaccg tcaatgccta caagaccatt gcccgagagg aagggttccg 300
gggcctctgg aaagggacct ctcccaatgt tgctcgtaat gccattgtca actgtgctga 360
gccggcgacc tatgacctca tcaaggatgc cctcctgaaa gccaacctca tgacagatga 420
cctcccttgc cacttcactt ctgcctttgg ggcaggcttc tgcaccactg tcctgcctc 480
ccctgtagac gtggtcaaga cgagatacat ga 512

<210> 1015
<211> 553
<212> DNA
<213> Homo sapien

<220>
<221> misc_feature
<222> (1)...(553)
<223> n = A,T,C or G

<400> 1015
ctgggcagga agattatgat cgcccagagg ccctctccta cccagatacc gatgttatac 60
tgatgtgttt ttccatcgac agccctgata gtccagaaaa catcccagaa aagtggaccc 120
cagaagtcaa gcattttctgt cccgacgtgc ccatcatcct ggttggaat aagaaggatc 180
ttcggaatga tgagcacaca aggcggggagc tagccaagat gaagcaggag cgggtgaaac 240
ctgaagaagg cagagatatg gcaaacagga ttggcgccct tgggtacatg gagtgtcag 300
caaagaccag agatggagtg agagaggttt ttgaaatggc tacgagagct gctctgcaag 360
ctagacgtgg gaagaaaaaa tctgggtgcc ttgtcttgtg aaaccttgct gcaagcacag 420
cccttatgcg gttaattttg aagtgtctgt tattaatctt agtgtatgat tactggcctt 480
tttcatttat ctataattta cctaagatta caaatcanga agtcatcttg ctaccagtat 540
ttagaagcca act 553

<210> 1016
<211> 431
<212> DNA
<213> Homo sapien

```

```
<210> 1017
<211> 490
<212> DNA
<213> Homo sapien
```

<400> 1017							
ctggaagaac	aaggcgaagt	tctggtggct	gtctgcgatg	aatgtgccct	tggctttggc		60
tgggtatgtc	acccgggtag	ttttgggtgc	aatgctctga	tccttatcca	cgggtggaaag		120
atcaacattt	gtgatgccaa	cttcagtggg	gatcttgact	ctgagctcta	cggatatttgc		180
aatataccgg	ttgtcacctt	caacttcgac	aaggaagtca	taataaccac	tggaaaattt		240
gacgttcatg	aaatttagtt	caaaaacatc	ccctacaggg	gtgaaggatg	tcttctggag		300
gacagtggct	ctggaagcaa	cagatttagc	atgttctagt	ttaacagtg	cctgagtcag		360
aggctgagac	agaacattgg	tgacttgcaa	ccgcaagata	gcctgttcat	gagtgtcgga		420
agcaganccc	tcangcacia	ccacaactgg	cacgtggtag	cgattatgcg	agagcacagg		480
cagacctcgg							490

<400>	1018						
ggagtaagct	gagtacaagt	accatagcag	cagagctgca	aaaggtcttg	ggacctatag		60
tcctaattgca	agataagggtc	atggggccta	aggccatggg	gcttgaggca	cccctagacc		120
ctgagccttc	agcatttaag	ggagggtgtc	ccccattct	cgataggcca	tggtacacag		180
atgggtctag	ccgaggtgct	ataactgctt	ggaccactgt	tgcagtccaa	cctagtactg		240
acactatatg	gtttgaaacc	cgggtgtggac	aaagtagcca	atgggctgaa	cttagagcag		300
tgtggatggg	gatcaccaag	gagggtgacac	tgatggtaat	ctgtatcaat	agctggggtg		360
tctaccaagg	cttaactttg	tggttaacta	cctggaaaat	acagaagttg	ctagtcggcc		420
accaacccat	ttggggtc aa	gccacgtggc	aagacctctg	ggaaatgggt	catcagaaac		480
aggtaaccgt	ttatcatgtg	tca					503

<400> 1019
cctgtgtatg gactagaggc ggggtgcacgg gtactgttcc tcacggcagt caagaggccc 60

aggtctctgtg	ggctccagct	ctgcatttcc	cggttctggg	gttggggctg	ggatgacttc	120
ctgttggact	tgtgtctggg	actggaactg	gaactgttcc	tcggagggcc	gaggagtcac	180
ctcttgataa	tcatagtagt	ctgggttgtc	gatctggtcg	ctatagtggg	tgtactggac	240
gtggtcaggg	aacggcggca	gcgggtccag	gtcatactgg	ccctgagcca	gcaagcctgc	300
aggcaggaat	agcaggaaga	ggtaggcagc	tctcatggca	acaaagag		348

<210> 1020

<211> 260

<212> DNA

<213> Homo sapien

<400> 1020

ccacacggcg	accgagggac	agatggggcc	ctgcgtccca	taggctgcct	gaaggtgggt	60
agggcggcct	gcggcatagt	ggggtggctg	tgggctccca	gcctggcccc	tgggaaccgt	120
gggagcacag	ggacaagcac	atggctatgg	aatgcagggg	gacccaagga	caagcgagtt	180
gcggggatct	ctactgtgac	catgcagaat	tgatcgcagt	ctgctgcgcc	accaccacct	240
catgttcccc	aggggaacag					260

<210> 1021

<211> 407

<212> DNA

<213> Homo sapien

<400> 1021

ccttatgact	ataacggccc	acgagaaaaa	tatggaatcg	ttgattacat	gatcgagcag	60
tccgggcctc	cctccaagga	gattctgacc	ctgaagcagg	tccaggagtt	cctgaaggat	120
ggagacgatg	tcatcatcat	cggggtcttt	aagggggaga	gtgaccacagc	ctaccagcaa	180
taccaggatg	ccgctaacaa	cctgagagaa	gattacaaat	ttcaccacac	tttcagcaca	240
gaaatagcaa	agttcttgaa	agtctcccag	gggcagttgg	ttgtaatgca	gcctgagaaa	300
ttccagtcca	agtatgagcc	ccggagccac	atgatggacg	tccagggctc	cacccaggac	360
tcggccatca	aggacttcgt	gctgaagtac	gccctgcccc	tggttgg		407

<210> 1022

<211> 140

<212> DNA

<213> Homo sapien

<400> 1022

ccaccccaga	gtgggagagg	ctgggagggt	gggaggctgt	ggagagaagt	gagcaagggtg	60
ctcttgaacc	tgtgtcatt	ttgcaatttt	atcagtaatt	tgacttagag	tttttacgaa	120
acctcttttg	ttgtccttgc					140

<210> 1023

<211> 280

<212> DNA

<213> Homo sapien

<400> 1023

ctggagggtgc	ctcagaagggt	gcattctgct	tcctgcaggg	gcttgaaaca	ccaaggcact	60
ccagggatcc	tggagtcaaa	gcagcagccc	cggttgttgc	actccttggg	ggtgacatgg	120
gggtagccgc	agtccaccct	gtccttggct	ggcacggcac	actggtttgc	agacaggccc	180
gcgtactcct	cagcagagct	ggaggacagc	aaggccagga	ccagccccag	catgcagagc	240
gctctggcag	ccatgaccac	cgtgggctcc	gggacgcagc			280

<210> 1024
 <211> 274
 <212> DNA
 <213> Homo sapien

<220>
 <221> misc_feature
 <222> (1)...(274)
 <223> n = A,T,C or G

<400> 1024
 cctggctgag caggcagagc accctgggac cccagggcag aaggaccct gccctccagt 60
 cccaagacc caggcccgtc tccactcata cagccacct acatgtgacg tcagccctga 120
 aaaggtaaca ggaaagtcca gaacaaaaaac aaaaccccaa aagtaaaaag gctacgtgta 180
 gcagagtaat accggaaacg ttatatacac aggcgggtgat ggccccctcg gaagtgtccg 240
 ggtcacttag ggggcactgc anaggtccct gtgg 274

<210> 1025
 <211> 446
 <212> DNA
 <213> Homo sapien

<220>
 <221> misc_feature
 <222> (1)...(446)
 <223> n = A,T,C or G

<400> 1025
 gcaaagagtg tactgtgctt gaggcagagc actcacacat aaatggctgt gtgtggaatt 60
 gcttgccaaa gaagtttcta gcctttccct tccccctaac tgcacaggg aagaattctt 120
 atctctagct tggtttccac atgaggtttt tctgagaagg gcttgggaca agaagtctgt 180
 catgttagtt aagcaggcaa gaaatcctac taatccagtt ttgtttgaaa gttgtttgtc 240
 cgtatgatatt tttaaaagtc aagtttaatt tcaaaaaacc tttttttct gagattactt 300
 ttggggtaat atttaaaatg agagacattt tgtaaccctg taaaatacat aggggaatata 360
 acattccagt gtatacaaag aaggcaaatt cttaaatcaa ataaagcgca ttataaaatc 420
 aaaaaanaaa naaaaaaaan aaaaaa 446

<210> 1026
 <211> 189
 <212> DNA
 <213> Homo sapien

<400> 1026
 ctgtgagaga gatgctcaat atgccccagg ctatgacaaa gtcaaggaca tctcagaggt 60
 ggtcacccct cggttccttt gtactggagg agtgagtccc tatgctgacc ccaatacttg 120
 cagaggtgat tctggcggcc ccttgatagt tcacaagaga agtcgtttca ttcaagttgg 180
 tgtaatcag 189

<210> 1027
 <211> 92
 <212> DNA
 <213> Homo sapien

<400> 1027

ccagaccctc cttagtagac gatctcggac cacaaaccaa ggagtctcgt ggccttggat 60
 tcccagaccc taggatggta tccctctgac ag 92

<210> 1028
 <211> 438
 <212> DNA
 <213> Homo sapien

<400> 1028
 ctgaaaagcc atcttttgc at tgttctcat cgcctcctt gctcgccgca gccgcctccg 60
 ccgcgcgcct cctccgccgc cgcggactcc ggcagcttta tcgccagagt ccctgaactc 120
 tcgctttctt tttaatcccc tgcctcggat caccggcgtg ccccaccatg tcagacgcag 180
 ccgtagacac cagctccgaa atcaccacca aggacttaaa ggagaagaag gaagttgtgg 240
 aagaggcaga aaatggaaga gacgcccctg ctaacgggaa tgctaataag gaaaatgggg 300
 agcaggaggc tgacaatgag gtagacgaag aagaggaaga aggtggggag gaagaggagg 360
 aggaagaaga aggtgatggt gaggaagagg atggagatga agatgaggaa gctgagtcag 420
 ctacggggca gcgggcag 438

<210> 1029
 <211> 330
 <212> DNA
 <213> Homo sapien

<400> 1029
 ccagccgcat gggagtggag gcagtcacgc ccttgctaga ggccaccccg gacacccag 60
 cttgcgtcgt gtcactgaac gggaaccacg ccgtgcgcct gccgctgatg gagtgcgtgc 120
 agatgactca ggatgtgcag aaggcgatgg acgagaggag atttcaagat gcggttcgac 180
 tccgagggag gagctttgcg ggcaacctga acacctacaa gcgacttgcc atcaagctgc 240
 cggatgatca gatcccaaag accaatcgca acgtagctgt catcaacgtg ggggcacccg 300
 cggctgggat gaacgcggcc gtacgctcag 330

<210> 1030
 <211> 228
 <212> DNA
 <213> Homo sapien

<400> 1030
 ctggagactc tgggccagga gaagctgaag ctggaggcgg agcttgccaa catgcagggg 60
 ctggtggagg acttcaagaa caagtatgag gatgagatca ataagcgtac agagatggag 120
 aacgaatttg tctcatcaa gaaggatgtg gatgaagctt acatgaacaa ggtagagctg 180
 gagtctcgcc tggaagggct gaccgacgag atcaacttcc tcaggcag 228

<210> 1031
 <211> 294
 <212> DNA
 <213> Homo sapien

<400> 1031
 ccacaaagcc attgtatgta gcttttagctc agcgcaaaga agagcgccag gctcacctca 60
 ctaaccagta tatgcagaga atggcaagtg tacgagctgt gcccacccct gtaatcaacc 120
 cctaccagcc agcacctcct tcaggttact tcatggcagc tatccacag actcagaacc 180
 gtgctgcata ctatcctcct agccaaattg ctcaactaag accaagtccc cgctggactg 240
 ctcagggtgc cagacctcat ccattccaaa atatgcccgg tgctatccgc ccag 294

<210> 1032
 <211> 278
 <212> DNA
 <213> Homo sapien

```
<400> 1032
ggaggtatta cagacagcac tgcactttgg agttgggcag ctacatcgag gacctctttg      60
tgggtccacag tgacctctcc agcattgtga tcttgataa ctccccaggg gcttacagga      120
gccatccaga caatgccatc cccatcaaat cctgggttcag tgacccagc gacacagccc      180
ttctcaacct gctcccaatg ctgggtgccc tcagggttcac cgctgatgtt cgttcctgtg      240
tgagccgaaa ccttcaccaa catcggtctt ggtgacgg      278
```

<210> 1033
 <211> 155
 <212> DNA
 <213> Homo sapien

<220>
 <221> misc_feature
 <222> (1)...(155)
 <223> n = A,T,C or G

```
<400> 1033
cgcgttcanc catgttnaaa ccgattgcat naacttcgaa accggccccg ccgcccggcg      60
ctggagaggg gcanngggag aagcagagag tttatcattc atctgtacac atagacgttt      120
cttcttttaa taacaccacg ggcgggagcc ccac      155
```

<210> 1034
 <211> 401
 <212> DNA
 <213> Homo sapien

```
<400> 1034
ctggaccagc acccattga cgggtacctc tcccacaccg agctggctcc actgcgtgct      60
ccccctatcc ccatggagca ttgcaccacc cgctttttcg agacctgtga cctggacaat      120
gacaagtata tcgccctgga tgagtgggcc ggctgcttcg gcatcaagca gaaggatata      180
gacaaggatc ttgtgatcta aatccactcc tcccacagta ccgattctc tctttaaccc      240
tccccttcgt gtttccccca atgtttaaaa tgtttgatg gtttggtgtt ctgcctggag      300
acaagggtgct aacatagatt taagtgaata cattaacggt gctaaaaatg aaaattctaa      360
cccaagacat gacattctta gctgtaactt aactattaag g      401
```

<210> 1035
 <211> 333
 <212> DNA
 <213> Homo sapien

```
<400> 1035
ctgagctggg ggttgaatth ctccaggcac tccctggaga gaggaccag tgacttgtcc      60
aagtttacac acgacactaa tctcccctgg ggaggaagcg ggaagccagc caggttgaac      120
tgtagcgagg cccccaggcc gccaggaatg gaccatgcag atcactgtca gtggaggga      180
gctgctgact gtgattaggt gctgggggtc tagcgtccag cgagcccgg gggcatcctg      240
gaggctctgc tccttagggc atggtagtca ccgcgaagcc gggcaccgct ccacagcatc      300
tcctagaagc agccggcaca ggaggggaagg tgg      333
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<210> 1036
 <211> 198
 <212> DNA
 <213> Homo sapien

<400> 1036	
ccaatgtaca tgggtggacta tgccggcctg aacgtgcagc tcccgggacc tcttaattac	60
tagacctcag tactgaatca ggacctcact cagaaagact aaaggaaatg taatttatgt	120
acaaaatgta tattcggata tgtatcgatg ccttttagtt tttccaatga tttttacact	180
atattcctgc caccaagg	198

<210> 1037
 <211> 289
 <212> DNA
 <213> Homo sapien

<400> 1037	
ctggagatga tcctcaacaa gccagggctc aagtacaagc ctgtctgcaa ccagggtggaa	60
tgtcatcctt acttcaacca gagaaaactg ctggatttct gcaagtcaaa agacattggt	120
ctggttgcct atagtgtctt gggatcccac cgagaagaac catgggtgga cccgaactcc	180
ccggtgctct tggaggaccc agtcctttgt gccttggcaa aaaagcacia gcgaaccca	240
gccctgattg ccctgcgcta ccagctacag cgtgggggtg tggtcctgg	289

<210> 1038
 <211> 368
 <212> DNA
 <213> Homo sapien

<400> 1038	
ccagacgtgg tggctcacac ctgcaatccc agcaccttag gaggccgagg caggaggatc	60
cttgagggtca ggagttcgag accagcctcg ccaacatggt gaaaccccat ttctactaaa	120
aatacaaaaa attagccaag tgtggtggca tatgcctgta atcccaacta ctcagaaggc	180
cgaggcagga gaattacttg aacgcaggag aatcactgca gcccaggagg cagaggttgc	240
agtgagccga gattgcacca ctgcactcca gcctgggtga cagagcaaga ctccatctca	300
gtaaataaat aaataaataa aaagcgctgc agtagctgtg gcctcaccct gaagtcagcg	360
ggccccagg	368

<210> 1039
 <211> 417
 <212> DNA
 <213> Homo sapien

<220>
 <221> misc_feature
 <222> (1)...(417)
 <223> n = A,T,C or G

<400> 1039	
ctgggcctat gctgggtcatg aacggctcctg gaaaatgact cccttccttc agtatctgca	60
tcctcatgaa gtcattcatt ttggagatcg tgtcttcact tttcttgggtg aagaaactgc	120
tggatggagt tgttgggtggc atctgaggag tccgaagatg gctctcaggg aaggttgtgc	180
tggcctctga aggatttggga agctgactct gttcctgggg tagctnnatg ctcttgggggt	240
cattgnttct cgggtttgnt tttttcttta tctggataaa actatgcatt tctgaaatca	300
gttttgacat ctggttcttt tttcctaagt cgaaagcaga aaagttggaa gcttatctcc	360

ttcttcacag ggggatattg tggacattgn nctgtcccca ctacatccat ttttcct 417

<210> 1040

<211> 409

<212> DNA

<213> Homo sapien

<400> 1040

ctgtccaatg	gcaacaggac	cctcactcca	ttcaatgtca	caagaaatga	cgcaagagcc	60
tatgtatgtg	gaatccagaa	ctcagtgagt	gcaaaccgca	gtgaccaggt	caccctggat	120
gtcctctatg	ggccggacac	ccccatcatt	tccccccag	actcgtctta	cctttcggga	180
gcgaacctca	acctctcctg	ccactcggcc	tctaaccocat	ccccgcagta	ttcttggcgt	240
atcaatggga	taccgcagca	acacacacaa	gttctcttta	tcgccaaaat	cacgccaaat	300
aataacggga	cctatgcctg	ttttgtctct	aacttggtta	ctggccgcaa	taattccata	360
gtcaagagca	tcacagtctc	tgcattctgga	acttctcctg	gtctctcag		409

<210> 1041

<211> 492

<212> DNA

<213> Homo sapien

<220>

<221> misc_feature

<222> (1)...(492)

<223> n = A,T,C or G

<400> 1041

cctcggtccc	acacctccgc	tgtgaccaca	gcctcaggtc	aagctgtgct	ggggccatcc	60
accttccttt	gccatttaga	agatggggct	tggagcttgg	caacacagaa	attgacatca	120
gccttataaa	accttggctg	aacctaccga	cctccaggag	aatttcagcc	aaaacaaaaa	180
agcaaataca	cagagggacc	ctggaaccag	aatccctccc	catgggaaaag	acgaaggcac	240
agagattcga	gccaagtttc	ccaacatggt	ggtgtttgca	gaaaagtccg	gtcacgtcac	300
acacagcaca	gaggcaagaa	gcgaaggcag	tggcattcac	aggactactt	tatattaaag	360
tttattacat	ttggaaaatc	tactgtacag	ggaaaaaccc	attggattaa	gtagagtttt	420
gccaaaagca	aaagactatc	actctttgga	aaatattcct	gattccagcc	cangggcccag	480
ggtggggcca	ca					492

<210> 1042

<211> 125

<212> DNA

<213> Homo sapien

<400> 1042

cctggctctg	atccagtgc	ccctctcacc	aaagaactcg	gtttaaccag	ggctctgtaa	60
gacctctccc	accagagac	ttgtgtggcc	tgggtgtggcc	tgtgtgtcgg	attccttcct	120
gtcag						125

<210> 1043

<211> 459

<212> DNA

<213> Homo sapien

<400> 1043

ccagcctgga	gataaggggtg	aaggtggtgc	ccccggactt	ccaggtatag	ctggacctcg	60
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<221> misc_feature
 <222> (1)...(412)
 <223> n = A,T,C or G

<400> 1047
 gtacaagctt tttttttttt tttttttttt tttgtttaat gcttgaactt ttttttgag 60
 agagaaat tttt agaaagacac aaggtacaca gagtaaaatg tttttctttt ttcaggacct 120
 tgaactgaat cttgcactgc tttggtttct atctaggaag ctcagcgaca gcagagtctg 180
 tanaggcggc cactgatttc acacaccccg gagagggact cacgggtagc acaacggccg 240
 gttcggcaat agcaggtggc tcttgccctga naacctgagg ttctaanaagc ananagtcca 300
 tttcctgcaa aggagatagc aaggtcctgg ttgtcttccc canactgctt ctgggttgta 360
 gcctcatcag ctctttcctg gagtgactca gcctgggctt gcagggccac ca 412

<210> 1048
 <211> 476
 <212> DNA
 <213> Homo sapien

<220>
 <221> misc_feature
 <222> (1)...(476)
 <223> n = A,T,C or G

<400> 1048
 taaaaaaagg aaaaagt tttt attacgaaac tagtttgtat aaaacaggggt tatacatatt 60
 tttgtaagtt tgtaataaaa cagtaagaaa aaaaggcagt aatagaaatc tccaaaaggc 120
 aacctatcaa aaccaactgg ctgccacttt gagtttggac agtagctgca taaactttgt 180
 tcttcttgaa cagtatttaa taacatcatt aatacattaa caacatttct ataaagtaag 240
 acacattgggt gctgaagtac aactggnggc ctcttgatct cacctatgag gagagttctt 300
 taaaaaacca catagggaaa attgcagttg taaggngaac tacncatcta aaatatgcan 360
 aggtaatagc attacatgtt aaaggtatca aggnatata cacattttta accatttggn 420
 acaaaacttn tataaaattt ntttctctct ctttctctct tatgcacaaa aaatat 476

<210> 1049
 <211> 274
 <212> DNA
 <213> Homo sapien

<400> 1049
 cctggctgag caggcagagc accctgggac cccagggcag aaggaccctt gccctccagt 60
 cccaagacc caggcccgct tccactcata cagccacct acatgtgacg tcagccctga 120
 aaaggtaaca ggaaagttca gaacaaaaac aaaaccccaa aagtaaaaag gctacgtgta 180
 gcagagtaat accggaaacg ttatatacac aggcggtgat ggccccctcg gaagtgtccg 240
 ggtcacttag ggggcactgc agaggtccct gtgg 274

<210> 1050
 <211> 472
 <212> DNA
 <213> Homo sapien

<400> 1050
 ctgcagcctg ggactgaccg ggaggctctg attatttacc caccacaggt aggttggtgtt 60
 ctgaatctca ggttcacagg ttaaggctac agcatcctca tctccacgg ggttgaggtt 120
 gttgctgggt atgaaggggt tgggtggctc tgcatagact gtgatcgctg tgactgtggt 180

cctattgagg	ccagtgtctg	agttatgggc	ttggcacgta	taggatccac	tattattcac	240
agtgatgttg	gggataaaga	gctcttgggt	ggattgctgg	aaagtcccat	tgacaaacca	300
agagtactgt	gcaggtgggt	tagaggctgc	gtggcaggag	aggttcagat	tttccctga	360
tctgtaagat	gtgttttagag	gggaaatggg	gggggcatcc	gggccataga	ggacattcag	420
gatgactgaa	tcactgcgcc	tggcactcac	tgggttctgg	gtttcacatt	tg	472

<210> 1051

<211> 249

<212> DNA

<213> Homo sapien

<400> 1051

ccaccaaccg	tggcatcacg	cgaatccggg	gcaccagcta	ccagagccct	cacggcatcc	60
ccatagacct	gctggaccgg	ctgcttatcg	tctccaccac	cccctacagc	gagaaagaca	120
cgaagcagat	cctccgcata	cgggtgcgag	aagaagatgt	ggagatgagt	gaggacgcct	180
acacggtgct	gacccgcata	gggctggaga	cgctactgcg	ctacgccata	cagctcatca	240
cagacctgc						249

<210> 1052

<211> 289

<212> DNA

<213> Homo sapien

<400> 1052

ccaggaccac	aacccccacg	tgtagctggt	agcgcagggc	aatcagggct	ggggttcgct	60
tgtgcttttt	tgccaaggca	caaaggactg	ggctctccaa	gagcaccggg	gagttcgggt	120
ccacccatcg	tttgtctcgt	tgagatccca	gagcactata	ggcaaccaga	acaatatctt	180
tgcacttgca	gaaatctagc	aatttactcc	ggttgaaata	cggatgacat	tctacctggt	240
tgcagacagg	cttgtaactg	agtctctggc	tggttgaggat	catctccag		289

<210> 1053

<211> 199

<212> DNA

<213> Homo sapien

<400> 1053

ccacgactgc	atgcccgcgc	ccgccagggt	atacctccgc	cggtgaccca	ggggctctgc	60
gacacaagga	gtctgcatgt	ctaagtgcta	gacatgctca	gctttgtgga	tacgcggact	120
ttgttgctgc	ttgcagtaac	cttatgccta	gcaacatgcc	aatctttaca	agaggaaacc	180
gtaagaaagg	gcccagccg					199

<210> 1054

<211> 224

<212> DNA

<213> Homo sapien

<400> 1054

tgcacctgt	gaagcaggag	acagatgctg	cattttcact	gttgtttgtc	ctctgttttt	60
gtagcatccc	cgggaacttc	cccatcagcc	aggggcttgt	ccccaccacc	cttcacctgg	120
ctttccagtt	ggctgagacg	ctgcttcata	ttcatctggg	tggcgttgta	ctcagccagg	180
aggcgtgcaa	acctggtctg	cagggcgctc	agggaggacc	ccag		224

<210> 1055

<211> 390

<212> DNA
<213> Homo sapien

<400> 1055
cctcttatta gggctctggt agcggcgggc gcggaacctt ggggtctgga cgcaacggcg 60
gcgggagcat gaacgccccct ccagccttcg agtcgttctt gctcttcgag ggcgagaaga 120
agatcaccat taacaaggac accaaggtac ccaatgcctg tttattcacc atcaacaaag 180
aagaccacac actgggaaac atcattaaat cacaactcct aaaagaccgc caagtgcctat 240
ttgtctggcta caaagtcccc cacccttgg agcacaagat catcatccga gtgcagacca 300
cgccggacta cagccccag gaagcctttg ccaacgccat caccgacctc atcagtgcgc 360
tgtccctgct ggaggagcgc tttcgggtgg 390

<210> 1056
<211> 450
<212> DNA
<213> Homo sapien

<220>
<221> misc_feature
<222> (1)...(450)
<223> n = A,T,C or G

<400> 1056
ccagcatcac cttttggtcc nnacactcca gggctgccag gagcaccagt gttaccgcga 60
ggacctgggg gccatcctt gcctggagaa ccgctgggac ctgggggtcc tgggttacca 120
ttactaccag gaggaccagg aagaccacga gcaccaggga agccagcagc accaggtcca 180
ccaggactgc caggttcacc tttgacacct tggggaccag gaggaccagn angtcagaa 240
cctccagggg gtcttgcaac tccaggaggg cctccttcac ctttctcacc cggagccccct 300
ctttctcctt taccaccagg ttcaccattc tgtccaggag caccagggaa accagcaggt 360
cctggagggc cagtttnacc tctctcacca nggctaccac gaggtccagc tatacctgga 420
agtcgggggg caccaccttc acccttacct 450

<210> 1057
<211> 337
<212> DNA
<213> Homo sapien

<400> 1057
tgagcgggcg cccggcaggt cctcgcctgg agggccccgg gcagcacagg gaggacgagc 60
ttgtccagca gagggtctgg cagaggtcc cgcagaggtt tgggcagggg gtctgacatc 120
cctggctcct gctctggctc tggctgccgg gatttgcaca ggcccaggtg catacagatg 180
ccgtttgagt caatctgggt ctggaagtag tcgatgacca gggggaagta gtcgtcaagc 240
acttggttgc actggggcat gagcagcttc aaggggagga cgttgactc ctgctccagg 300
aacttctca tcgtgtcctg gaaaatggcc tccttgg 337

<210> 1058
<211> 237
<212> DNA
<213> Homo sapien

<400> 1058
ctgggggactg ggaatgctag catatggtat ctcaagttgg ctctcagaac taaacgggga 60
taagggccta gaatggaaga gggaaccagc cagacctca gtccttcctg tcctggactg 120
ggagccacag atgtccctgt gatctgtcac tgccctgatc tgggtcttca gccattaaa 180

ctcagtgtca tcttcagtca ccaacggggg tcttggtgtc cttccaaacc cctttgg 237

<210> 1059

<211> 210

<212> DNA

<213> Homo sapien

<220>

<221> misc_feature

<222> (1)...(210)

<223> n = A,T,C or G

<400> 1059

agcccatccc	cccggtccc	tcctagtctg	ccctgcgtcc	tctgtccccg	ggtttcagag	60
acaacttccc	aaagcacaaa	gcagtttttc	cccctagggg	tgggaggaag	caaaagactc	120
tgtacctact	ttgtatgtgt	ataataattt	gagatgtttt	taattattnn	gattgctgga	180
ataaagcatg	tggaatgac	ccaaaaaaaa				210

<210> 1060

<211> 564

<212> DNA

<213> Homo sapien

<400> 1060

ctggccacag	agcccagcaa	gtccttcctg	ggagagaaga	gttagggctg	atactgaagg	60
tctctttcac	atctgggcac	acgtctgcct	tcaggctgta	agaatttcat	ttgtcgattg	120
ttaaataaaa	ccaggagaaa	gcaatgcagg	tctctgggaa	tctcatccct	tccataagga	180
aaatgctctg	ccaattcaag	tttcattcag	tcaggaagac	agaaggattt	aaggcttcgg	240
tgacaattat	aatcctctga	gaaattattt	ccccttaaag	tcaagataag	ataatagtgt	300
ttactgtact	ttctcttgac	tcttgaaatc	cctggtattg	ggtgtaggca	acttgcacct	360
gcaatgaagt	ccgcaggaga	ggaaggtctc	tcctcccccg	aaagctatcc	cagggtcacat	420
gcgtggcgaa	tgcccaactga	acctcggctc	tcatggaagc	aggaaagaca	ccgagattca	480
agccttctag	taggttgagg	acgctgtgct	catggcatct	tcggagattt	tggtactggc	540
aggggtggat	gcttgcaaaa	tact				564

<210> 1061

<211> 267

<212> DNA

<213> Homo sapien

<400> 1061

cctatggagg	tgccatgat	gtcatgagct	ctaagcacct	ttgtggtgat	accaactatg	60
cctggccac	cgagagatt	gcggtcatgg	gagcaaaggg	cgctgtggag	atcatcttca	120
aagggcatga	gaatgtggaa	gctgctcagg	cagagtacat	cgagaagttt	gccaaacctt	180
tcctgcagc	agtgcgagg	tttgtggatg	acatcatcca	accttcttcc	acacgtgccc	240
gaatctgctg	tgacctggat	gtcttgg				267

<210> 1062

<211> 603

<212> DNA

<213> Homo sapien

<220>

<221> misc_feature

<223> n = A, T, C or G

ctggtcatct	tgctcatgtga	agaccatctt	cctacagagt	ctaggtctggc	cgtcgttgaa	60
gtcctcacca	gtactacacc	acttttcttc	accaaccccc	atcctattct	tgagttgcag	120
gatacacttg	ctctctggaa	gtgtgtcctt	acccttctgc	agagtgagga	gcaagctggt	180
agagatgcag	ccacggaaac	cgtgacaact	gccatgtcac	aagaaaatac	ctgccagtca	240
acagagtttg	ccttctgcca	ggtggatgcc	tccatcgctc	tggccctggc	cctggccgtc	300
ctgtgtgatc	tgctccagca	gtgggaccag	ttggcccttg	gactgcccat	cctgctggga	360
tggctgttgg	gagagagtga	tgacctcgtg	gcctgtgtgg	agagcatgca	tcagggtggaa	420
gaagactacc	tgtttgaaaa	agcagaagtc	aacttttggg	ccgagaccct	gatctttgtg	480
aaatacctct	gcaagcacct	cttctgtctc	ctctcaaaag	tccggtctggc	gtncoccaaag	540
ccctgagatg	ctctgtcacc	ttcaaaggat	ggtgtcagag	cagtgccacc	tnctgtctca	600
qtt						603

<211> 222

<213> Homo sapien

ccatcgtgga	tcactgagat	gcagtggcgg	tccccgtagc	tggccccgtg	catgccaccc	60
tggaagatga	tgaagggcaa	cccctgccta	gtggtcagcc	ggaggattct	ggtaatcgt	120
ctgcaaggaa	agggaccgta	aggcacgagg	ctgcggaggg	gctctggttg	ctgggcttcg	180
ctggacacgg	gccactggca	gtagctgccg	tcagagtgc	ag		222

<211> 72

<213> Homo sapien

<221> misc feature

<223> n = A, T, C or G

gatgatcaat atnnactgga acacatgcat gcttttggaa tgtataatta cctgcactgt 60
gattcatggt at 72

<211> 251

<213> Homo sapien

gtggccgtga	tggatagcga	caccacaggc	aagctgggct	ttgaggaatt	caagtacttg	60
tgaacaata	tcaaaagggtg	gcaggccata	tacaaacagt	tcgacactga	ccgatcaggg	120
accatttgcg	gtagtgaact	cccagggtgcc	tttgaggcag	caggggttcca	cctgaatgag	180
catctctata	acatgatcat	ccgacgctac	tcagatgaaa	gtgggaacat	ggattttgac	240
aacttcatca	g					251

<210> 1066

<211> 289
 <212> DNA
 <213> Homo sapien

<400> 1066
 ctggagatga tcctcaacaa gccagggctc aagtacaagc ctgtctgcaa ccaggtggaa 60
 tgtcatcctt acttcaacca gagaaaactg ctggatttct gcaagtcaaa agacattgtt 120
 ctggttgctt atagtgtctt gggatccac cgagaagaac catgggtgga cccgaactcc 180
 ccagtgtctt tggaggaccc agtcctttgt gccttggcaa aaaagcacia gcgaacccca 240
 gccctgattg cctcgcgcta ccagctacag cgtgggggtg tggctctgg 289

<210> 1067
 <211> 301
 <212> DNA
 <213> Homo sapien

<400> 1067
 ctgtagttag ctgaagtcgc taaacaggac ggatttaagt agaggtgata tgtccagtca 60
 ccggcataga gacgtcctct gcgtcaccat ccacacacag ggcttctggg agacatcagg 120
 caaagctctc catgttaata ttcatctgaa tatggataat taggggtggc agcaaaacta 180
 tcaactgtta aatagtggag atttctgtct aggccatcta tggctttcat gtctccgca 240
 gtcaactgga actcaaaaac ctgcacgttc tgtctgatgc gctgctcatt gtagctcttg 300
 g 301

<210> 1068
 <211> 255
 <212> DNA
 <213> Homo sapien

<400> 1068
 ccagcagttc ctctttgcct tatatttggt gtacgcccgg ccagccttca agatggggtt 60
 gtcaattcgg ccacctccag ccaccacacc aaccacagct ctggtggctg aggagataac 120
 cttcttgagg ccggagggca gttcacacg ggtcttcttg gtctcagggt tgtgggagat 180
 aacggtggca tagttccctg atgcccgggc cagcttgcca cggctctccag gcttctctc 240
 caggcagcac acgat 255

<210> 1069
 <211> 77
 <212> DNA
 <213> Homo sapien

<400> 1069
 ctggacaggc tccagcaccg gcccaaacac gccagacct cggcaggcac cacctgggtc 60
 tcccacccag aaagtcc 77

<210> 1070
 <211> 163
 <212> DNA
 <213> Homo sapien

<220>
 <221> misc_feature
 <222> (1)...(163)
 <223> n = A,T,C or G

<400> 1070
 ctgctgggat gncgtccaag tttttcagcc ataaggtagc gaaatctagc agaatccaga 60
 ttacatccac ttccaatcac gcggtgtttg ggtaatccac ctagtttnna ggtaacatac 120
 gtaagaatgt ccactgngtt ggaaacmnca attatgatgc aat 163

<210> 1071
 <211> 246
 <212> DNA
 <213> Homo sapien

<220>
 <221> misc_feature
 <222> (1)...(246)
 <223> n = A,T,C or G

<400> 1071
 ctgaccggac cggncatgcc cgtccggaac gtctataaga aggagaaagc tcgagtcac 60
 actgaggaag agaagaattt caaagccttc gctagtctcc gtatggcccg tgccaacgcc 120
 cggctcttcg gcatacgggc aaaaagagcc aaggaagccg cagaacagga tgttgaaaag 180
 aaaaaataaa gccctcctgg ggacttgga tcagtcggca gacaaaaaaa aaaaaaaaaa 240
 aacaaa 246

<210> 1072
 <211> 224
 <212> DNA
 <213> Homo sapien

<220>
 <221> misc_feature
 <222> (1)...(224)
 <223> n = A,T,C or G

<400> 1072
 ctgccctgac agagcgctcc ttgatgggca tggactggaa aggatcccag gaatacaaga 60
 aggcagaaaa aaaagtittg aagatcttta aatctgacag tgaagtggct ggttacatcc 120
 ggcaagcggg tgacttccat cangtaatta ttcgaggtgg aggacatatt ttaccctatg 180
 accagcctct gagagctttt gacatgatta atcgattcat ttat 224

<210> 1073
 <211> 301
 <212> DNA
 <213> Homo sapien

<400> 1073
 ctgtagttag ctgaagtcgc taaacaggac ggatttaagt agaggtgata tgtccagtca 60
 ccggcataga gacgtcctct gogtcacat ccacacacag ggcttctggg agacatcagg 120
 caaagctctc catgttaata ttcattctgaa tatggataat taggggtggc agcaaaacta 180
 tcactgttaa aatagtggag atttctgtct aggccatcta tggctttcat gtccctctgca 240
 gtcaactgga actcaaaaac ctgcacgttc tgtctgatgc gctgctcatt gtagctcttg 300
 g 301

<210> 1074
 <211> 132

<212> DNA
<213> Homo sapien

<220>
<221> misc_feature
<222> (1)...(132)
<223> n = A,T,C or G

<400> 1074	
caagcttttt tttttttttt tttttttttt ttcgctcaaa nactttnttt tattantaca	60
tgggctggna ttgatggnaa gggacaaatg tanttggcaa ccatgggttag catcggatgc	120
ccatcccaat gg	132

<210> 1075
<211> 301
<212> DNA
<213> Homo sapien

<400> 1075	
ctgtagttga ctgaagtcgc taaacaggac ggattttaagt agaggtgata tgtccagtca	60
ccggcataga gacgtcctct gcgtcaccat ccacacacag ggcttctggg agacatcagg	120
caaagctctc catgttaata ttcattctgaa tatggataat taggggtggc agcaaaaacta	180
tcactgttaa aatagtgagg atttctgtct aggccatcta tggctttcat gtccctctgca	240
gtcaactgga actcaaaaac ctgcacgttc tgtctgatgc gctgctcatt gtagctcttg	300
g	301

<210> 1076
<211> 436
<212> DNA
<213> Homo sapien

<400> 1076	
ctgctgggat gaatgccaag tttttcagcc ataaggtagc gaaatctagc agaatccaga	60
ttacatccac ttccaatcac gcgggtgtttg ggtaatccac ctagtttcca ggtaacatac	120
gtaagaatgt ccactgggtt ggaaaccaca attatgatgc aatcaggact gtacttgacg	180
atctgaggaa taatgaattt gaagacatta acatttctct gcaccagatt gagccgactc	240
tcccttctct gctgacggac tcctgcagtt actactacaa tcttagaatt ggcggtcaca	300
gaataatctt tatctgccac aatttttaggt gtctgaagaa ataagctccc atgctgcaga	360
tccatcattt ctctttaaag cttatcttcc aaaacatcca caagagcaag ttcattcagcc	420
agagactttc ccagaa	436

<210> 1077
<211> 256
<212> DNA
<213> Homo sapien

<400> 1077	
ctgaagatta ataggaaaca gtgaaaaagc aacgtcctgt gatcagtaac tttaaagaca	60
agcttggttc tctctttctg gcaactactga cattcccacc attctagctt ccgaattctg	120
gaaaaagaga agatgattaa caaaaataga gaatgtagaa acttctgggt ttgtgcctac	180
aggattggca ccagaccctc agtgcctact tgctccatct acaaggcagc accctctcca	240
gaggcagcca gggagg	256

<210> 1078

<211> 202
 <212> DNA
 <213> Homo sapien

<220>
 <221> misc_feature
 <222> (1)...(202)
 <223> n = A,T,C or G

<400> 1078
 ctgtgctncn caaccagatc catgtnaagt gccccgcccc gagaagggag ccaggggggag 60
 ctgactncag ncaacancca gtgnccggat gancaccaac atgtgagggg tgaaccttgg 120
 cctccangac atntgcaccc cctnccccacc tccaoggacc tgggacctcc aggcggctca 180
 gtgctgcctg cggcccagct aa 202

<210> 1079
 <211> 170
 <212> DNA
 <213> Homo sapien

<400> 1079
 gcgcttctcg ggcaccgtca ggcttaagtc cactccccgc cctaagttct ctgtgtgtgt 60
 cctgggggac cagcagcact gtgacgaggc taaggccgtg gatatcccc acatggacat 120
 cgaggcgctg aaaaaactca acaagaataa aaaactggtc aagaagctgg 170

<210> 1080
 <211> 494
 <212> DNA
 <213> Homo sapien

<400> 1080
 cctgcggcaa agagatgcgc ttattgagaa acatggctta gttataatcc ccgatggcac 60
 tcccaatggt gatgtcagtc atgaaccagt ggctggagcc atcactgttg tgtctcagga 120
 agctgtctcag gtcttggagt cagcaggaga agggccatta gatgtaaggc tacgaaaact 180
 tgcctggagag aaggaagaac tactgtcaca gattagaaaa ctgaagcttc agttagagga 240
 ggaacgacag aaatgctcca ggaatgatgg cacagtgggt gacctggcag gactgcagaa 300
 tggctcagac ttgcagttca tcgaaatgca gagagatgcc aatagacaaa ttagcgaata 360
 caaatttaag ctttcaaaag cagaacagga tataactacc ttggagcaaa gtattagccg 420
 gcttgagggg caggttctga gatataaaac tgctgctgag aatgctgagg aaagtgaag 480
 atgaattgaa agca 494

<210> 1081
 <211> 123
 <212> DNA
 <213> Homo sapien

<400> 1081
 ctgctgctat taagttgcaa gctctacagc tagctacatg actgatggat cagtttgaga 60
 tttgttcctt tgtcaaaaagt ttaactctga tagaaggttg gcctcacatt ctgatgtttg 120
 gac 123

<210> 1082
 <211> 297
 <212> DNA

<212> DNA

<213> Homo sapien

<400> 1086
 cctcagaggt ttctccacag tcctcttctg ggcaaattct tgtttcttca catgccggac 60
 tagcttaaga ccaatgcagt agcttatttc caagccttgc aaagtatata atatctaaga 120
 ggaaagggtt tgtcatcca gcgttgtcca ctttgtgggg ctttgtaggt agacggagcc 180
 acactacagg caggggtatga gcagagggat gtatggagtg tgggtgactc tgagcctcac 240
 tgccgctgca aggtggggaa actgtaagtg aaccctgtg ggtgcggggg agggatatccg 300
 gtgcgcaggg aggtgg 316

<210> 1087

<211> 329

<212> DNA

<213> Homo sapien

<400> 1087
 cctgcagggg atgggacctt ccagaagtgg gcgtctgtgg tggcgccttc tggacaggag 60
 cagagataca cctgccatgt gcagcatgag ggtctgcccc agccctcac cctgagatgg 120
 gagcgtctt cccagccac catcccatc gtgggcatca ttgctggcct ggttctctt 180
 ggagctgtga tcgctggagc tgtggtcgct gctgtgatgt ggaggaggaa gagctcagat 240
 agaaaaggag ggagctactc tcaggctgca agcagtgaca gtgcccaggg ctctgatatg 300
 tctccacag cttgtaaagt gtgagacag 329

<210> 1088

<211> 342

<212> DNA

<213> Homo sapien

<400> 1088
 ccactcactg ctgggaccca ggcacctccc ttctccatcc tctctggatt gtcagtaatg 60
 tcctggaaca gaagcctgtg ggatggcctt gggcacggag aagccctggg gtcagtgtcg 120
 tgcacggatg gcggcagtgt tgaaccacagg aggctgaacc cggcccacca cggaagatga 180
 gtgcatggca accgcctgcc ttcacgtcgc tccacttggg aacccaagg tctgggctgt 240
 tctaggtatt gcttcacgtg cccagcaag cccttaacaa gagggcctgg ttccctgaag 300
 aaccaatccc aggaaggggc cttgatccct ccgccttgct ga 342

<210> 1089

<211> 51

<212> DNA

<213> Homo sapien

<220>

<221> misc_feature

<222> (1)...(51)

<223> n = A,T,C or G

<400> 1089

ccttggtgtc agtctccncg ctcttcttgc cactgttgag ggtggagatg t 51

<210> 1090

<211> 515

<212> DNA

<213> Homo sapien

```
<210> 1091
<211> 277
<212> DNA
<213> Homo sapien
```

```
<210> 1092
<211> 368
<212> DNA
<213> Homo sapien
```

```
<210> 1093
<211> 459
<212> DNA
<213> Homo sapien
```

```
<210> 1094
<211> 610
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<212> DNA
<213> Homo sapien

<220>
<221> misc_feature
<222> (1)...(610)
<223> n = A,T,C or G

<400> 1094
ccatgcaaaa ggaggtggtg cactcagtgc agtcgctgcc acaaaaagtc cgattatattt 60
cattggtaca ggggaacata tagatgactt tgaacctttc aaaacacagc cttttatttag 120
caaacttctt ggtatgggcg acattgaagg actgatagat aaagtcaacg agttgaagtt 180
ggatgacaat gaagcactta tagagaagtt gaaacatggt cagtttacgt tgcgagacat 240
gtatgagcaa tttcaaaata tcatgaaaat gggccccttc agtcagatct tggggatgat 300
ccctggtttt gggacagatt ttatgagcaa aggaaatgaa caggagtcaa tggcaaggct 360
aaagaaatta atgacaataa tggatagtat gaatgatcaa gaactagaca gtacggatgg 420
tgccaaagtt tttagttaaac aaccaggaag aatccaaaga gtagcaagag gatcgggtgt 480
atcaacaaga gatgttcgag aacttttgac acaatatacc aagtttgcac agatggtaaa 540
aaagatggga ggtatcaaag gacttttcaa aggtgggcga catgtctaan aatgtgagcc 600
agtcacagat 610

<210> 1095
<211> 232
<212> DNA
<213> Homo sapien

<400> 1095
ccttattttct cttgtccttt cgtacagga ggaatttgaa gtagatagaa accgacctgg 60
attactccgg tctgaactca gatcacgtag gactttaatc gttgaacaaa cgaaccttta 120
atagcggctg caccatcggg atgtcctgat ccaacatcga ggtcgtaaac cctattgttg 180
atatggactc tagaatagga ttgcgctggt atccctaggg taacttgttc cg 232

<210> 1096
<211> 377
<212> DNA
<213> Homo sapien

<400> 1096
ccacgctcat ggaaaccacc caaggacagc cagagtcac attccctggc aagctgggtg 60
tattcttcca aaagtttccc acccagtggt tcagacaggt gtagcgtctc tgcaggggtcc 120
cgtgcaatga agtcaaagtc ctcaggcagg aaagccaggc aggcacccag tctggcagcc 180
tctcgaacca gccacgcaca tgttttaaag ttctgttgct tgtctggcgt cgatgttacc 240
tggcacacag ccaccagggg cagttcgcag gaggaagagg agatagccat ggctctgggc 300
ctgggctgag cacaaggtac tgagagttga ggtatccgga gtccaggaca cagaaggagc 360
aggaatctgt gaggagg 377

<210> 1097
<211> 311
<212> DNA
<213> Homo sapien

<400> 1097
ccacgccatg gggctggagc actcccaaga ccctggggcc ctgatggcac ccatttacac 60
ctacaccaag aacttccgtc tgtcccagga tgacatcaag ggcattcagg agctctatgg 120

ggcctctcct gacattgacc ttggcaccgg cccaccccc acactggggc ctgtcactcc 180
 tgagatctgc aaacaggaca ttgtatttga tggcatcgct cagatccgtg gtgagatctt 240
 cttcttcaag gaccggttca tttggcggac tgtgacgcca cgtgacaagc ccatggggcc 300
 cctgctggtg g 311

<210> 1098

<211> 404

<212> DNA

<213> Homo sapien

<400> 1098

ccacccacgc ttaggttccc atcacactga tgactccggg tttggcgagc acaggagcgc 60
 aaaccttttc acattctttc tgtgatccaa atttgttttc gtttccacca caacctccat 120
 accagaatct tgcacagctt ttggtgtttg gatcatagta ccattttaat atgaaatccc 180
 tgcaagtccc ttcgtctttc ggcaacttgc atatatctgt ttcagtgaga gccaatgggt 240
 ctgtgctcac cattagattg atggttgaac tagaagctga ccttgctggc tgtggagggt 300
 ggggctgaga tttctttgta ctgaaacttc cgtggtagggt ggctctgacc tgagacctca 360
 ggtagcagac cacagccaca tggatatgtct gcccagcgag cagg 404

<210> 1099

<211> 442

<212> DNA

<213> Homo sapien

<400> 1099

ccatgggatg gctcttctga ccattggggg ccaggccagg ccaggccagg cttagggtag 60
 caaggaccag gccaaagggg cagggcctcc tttggagggg ttgaggggta catcctcggc 120
 tgggtgtttgc atccaggggt ccagcaggat ctcttccagt gagggtcggg aagaaggttt 180
 gggggccagg caccggcgga ttagggcaca gcagtctggg gagacatggg ctgggaagtg 240
 gagctcagct tccagaatct cctggtccct ctcaaagggg atgtccccac acaccatgtc 300
 atagaggagg atgccagtg accagacagt ggccgggagt gcatgggtact ggtgtcgaga 360
 gatccactct ggggggctgt acacccttgt cccatcaaag tcagtgtagg gttcatcatg 420
 aagcagggca ccaggaacca aa 442

<210> 1100

<211> 191

<212> DNA

<213> Homo sapien

<400> 1100

ccacgaaaat caatgagaag ccacaggtga tcgcggacta tgagagcgga cgggccatac 60
 ccaataacca ggtgcttggc aaaatcgagc gggccattgg cctcaagctc cggggaaagg 120
 acattggaaa gcccatcgag aaggggccta gggcgaaatg aacacaaagc ctcgaaatca 180
 gtgcgctcca g 191

<210> 1101

<211> 178

<212> DNA

<213> Homo sapien

<400> 1101

cgggtacttt ggtggacatg aaggaactgg gcatatggga gccattggct gtgaagctgc 60
 agacttataa gacagcagtg gagacggcag ttctgtact gcgaattgat gacatcgttt 120
 caggccacaa aaagaaaggc gatgaccaga gccggcaagg cggggctcct gatgctgg 178

<400> 1105
 ctggggccac tgcgggcac atgattggag tgctgggttg ggttgctctg atatagcagc 60
 cctgggtgtag tttcttcatt tcaggaagac tgacagttgt tttgcttctt ccttaaagca 120
 tttgcaacag ctacagtcta aaattgcttc tttaccaagg atatttacgg aaaagactct 180
 gaccagagat cgagaccatc ctagccaaca tcgtgaaacc ccatctctac taaaaatata 240
 gaaattagct ggacatggtg gcatgtgcct gtaatccag ctactcagga ggctgaggca 300
 ggagaactgc ttgaacaggg acccgggagg cggagattgg agtgagccga gatcgcgcca 360
 ctgcactcca gtctgggcta cacagtgaga ctctgtctca agaaaaataa acagaagaat 420
 tgggggttg gggtgggaaa cagtgtttcc aggcagagag aacagcacgt acaaaggaga 480
 ctgttgggag gggttaaatga aataattcat gtaagggtact tagtaccaca catgaatttc 540
 acaagcagca g 551

<210> 1106
 <211> 280
 <212> DNA
 <213> Homo sapien

<400> 1106
 ctgctcttca cacagggttc tggggaaaaac aaggaagaga tcatcaatta tgaatttgac 60
 accaaggacc tgggtgtgcct gggcctgagc agcatcggtg gcgtctggta cctgctgagg 120
 aagcactgga ttgccaacaa cctttttggc ctggccttct cccttaatgg agtagggctc 180
 ctgcacctca acaatgtcag cactggctgc atcctgctgg gcggactctt catctacgat 240
 gtcttctggg tatttggcac caatgtgatg gtgacagtgg 280

<210> 1107
 <211> 570
 <212> DNA
 <213> Homo sapien

<400> 1107
 ctgattagt tctaaggaat ggtccaatac tgttgccctt ttccttgact attacactgc 60
 ctggaggata gcagagaagc ctgtctgtac ttcattcaaa aagccaaaat agagagtata 120
 cagtccatga gaattcctct atttgttcag atctcataga tgacccccag gtattgtctt 180
 ttgacatcca gcagtccaag gtattgagac atattactgg aagtaagaaa tattactata 240
 attgagaact acagctttta agattgtact tttatcttaa aagggtggta gttttcccta 300
 aaatacttat tatgtaaggg tcattagaca aatgtcttga agtagacatg gaatttatga 360
 atggttcttt atcattttct tttccctttt ttggcatcct ggcttgctc cagttttagg 420
 tccttttagt tgcttctgta agcaacggga acacctgctg agggggctct ttcctcatg 480
 tatacttcaa gtaagatcaa gaatcttttg tgaaattata gaaatttact atgtaaatgc 540
 ttgatggaat tttttcctgc tagtgtagct 570

<210> 1108
 <211> 386
 <212> DNA
 <213> Homo sapien

<400> 1108
 ctgttctctg ggtgacactg tataaacacg atgaccctgc cttgacttta gttgctggtc 60
 ttacatcaaa taagcccaca gacaaactcc gtgccctgcc tctgtgggta tctttacaat 120
 acttgggact tgatggggtt gtggagagga tcaagcatgc ctgtcaactg agtcaacggg 180
 tgcaggaaaag tttgaagaaa gtgaattaca tcaaaatctt ggtggaagat gagctcagct 240
 cccagtggt ggtgttcaga tttttccagg aattaccagg ctgagatccg gtgtttaaag 300
 ccgtcccagt gcccaacatg acaccttcag gagtcggccg ggagaggcac tcgtgtgacg 360
 cgctgaatcg ctggctggga gaacag 386

```
<210> 1113
<211> 646
<212> DNA
<213> Homo sapien
```

<223> n = A, T, C or G

<213> Homo sapien

<213> Homo sapien

<213> Homo sapien

```
<400> 1116
ccttattttct cttgtccttt cgtacagggg ggaatttgaa gtagatagaa accgacctgg      60
attactccgg tctgaactca gatacgtag gactttaatc gttgaacaaa cgaaccttta      120
```

```
<210> 1117
<211> 370
<212> DNA
<213> Homo sapien
```

```
<210> 1118
<211> 494
<212> DNA
<213> Homo sapien
```

```
<210> 1119
<211> 407
<212> DNA
<213> Homo sapien
```

```
<210> 1120
<211> 548
<212> DNA
<213> Homo sapien
```

```
<400> 1123
ccaattgaaa caaacagttc tgagaccgtt cttccactac tgattaagag tgggggtggca      60
```

ggtattaggg ataatattca tttagccttc tgagctttct gggcagactt ggtgaccttg 120
 ccagctccag cagccttctt gtccactgct ttgatgacac ccaccgcaac tgtctgtctc 180
 atatcacgaa cagcaaagcg acccaaaggt ggatagtctg agaagctctc aacacacatg 240
 ggcttgccag gaaccatatc aacaatggca gcatcaccag acttcaagaa tttagggcca 300
 tcttcagct ttttaccaga acggcgatca atcttttctc tcagctcagc aaacttgcac 360
 gcaatgtgag ccgtgtggca atccaatata ggggcatagc cggcgcttat ttggcctgga 420
 tggttcagga taatcacctg agcagtgaag ccag 454

<210> 1124

<211> 219

<212> DNA

<213> Homo sapien

<400> 1124

cctgctccag agcacggctg accatttctg ctccgggata tcagctcccg ttccccaage 60
 acactcctag ctgctccagt ctcagcctgg gcagcttccc cctgcctttt gcacgtttgc 120
 atccccagca tttcctgagt tataaggcca caggagtggg tagctgtttt cacctaaagg 180
 aaaagccac ccgaatcttg tagaaatatt caaactaat 219

<210> 1125

<211> 246

<212> DNA

<213> Homo sapien

<400> 1125

ccagagctgg gcccaagctg cgctggaatc gcagcaggag aggggagtg gctggttctt 60
 cccaccactt cccaggctct gacagccgag actcatttcc aaggcacagc agctttctaa 120
 agggactgag tttggactgg gttttggacc tccaggggct ggagcttcat cacctgggca 180
 gtgtcttttc tcagagagca ggtttcttta tagtttgga ataaatgggt cacggttcaa 240
 aagaaa 246

<210> 1126

<211> 227

<212> DNA

<213> Homo sapien

<400> 1126

ccattgttcc cgtgcacga agcttgcagg cagcttcagg tcctcggtaa acataactct 60
 ctgggggtggc ttggggccac ccaggaaggt accacatagc ctcttcaagt agctcatgtc 120
 cacgttgtag aagttgtggc cggcctgcca cgtggtattc cgtttggtga catagttgac 180
 cagctcatcc gacaggggat ggaaagaggg cctgctccgg gcattgg 227

<210> 1127

<211> 377

<212> DNA

<213> Homo sapien

<400> 1127

cctgccgtcg atgccaggga ggccgacagg accttctttt ccagcggggc cgatatttcc 60
 aggggaacca ggaagacctc tgggtcccat gagaccagga tccccagggc gaccagcatc 120
 tccattaggt cctcggactc cagcagggcc acttgcacca cgactaccag gagggcccat 180
 gacgccagct ctgccatcag ctccaggaag accacagaga ccaggactac ctctcagccc 240
 aggaggtcct ggagggcccg cagatccagc ttccccatta gggcctctct ttccttcttc 300
 accactggga ccaggaggac cttggggccc agcagagccg ggctcaccct tgttaccgct 360

377

<213> Homo sapien

<213> Homo sapien

<213> Homo sapien

<213> Homo sapien

<211> 304

<213> Homo sapien

ccaccccgga	gatgacacga	ggctcacatg	actctagaca	cttggtggaa	agtgaggcga	60
gaaaaacaat	gacttggggc	aattacacga	ctgcaaagct	agagctgcca	acagggctcc	120
agggagcttg	gcttctgtag	aagttctaag	gaagcggtag	gaactccacg	gcggtggggc	180
gctaactagc	agggaccctt	gcaagtgttg	gtcggggggc	tcgagctgcc	tgagctgaca	240
cgaggggagg	ggtctgtgta	gccaacaggt	gaccgaaggg	cttgcttgcc	cacagcttac	300
ttgg						304

<211> 224

<213> Homo sapien

ctgacat	ctatagtaga	tatggaggag	gtccaagact	aactgtgaaa	gccctgtgta	60
aggaatgtgt	agtagaacgt	tgtcgcatat	tgcgtctgaa	gaaccaacta	aatgaagatt	120
ataaaactgt	taataatctg	ctgaaagcag	cagtaaaggg	cagcgatgga	ttttgggtgg	180
qaaagtccctc	cttgccggagt	tggcgccagc	tagctcttga	acag		224

<211> 250

<213> Homo sapien

cctactctgc	tgaggtggcg	cttcctgcta	agggcccttc	tctgcccttt	ctgccctcct	60
tccactccca	catgctgagc	cgccacaaag	accaaagaag	tgatggcttt	tctctgtccc	120
ctcgtgctct	gaggggagag	gggtgggtct	cctgagccac	tcagatggga	aagtccctta	180
ctcggccctt	ccctcccag	cagccccaag	ctttacactg	gatgcagcga	tcaaccacc	240
actcaccagg						250

<211> 315

<213> Homo sapien

ccaatgggct	ttgctgtagc	ttgctgaaat	caccaagcag	gagagattta	accagaggcg	60
atgtgtccag	tcaccagcat	agagccatcc	tctgtgtcac	catccacacg	cagggccttc	120
tggtagacct	catgcaatgc	cctccatggt	aatattcatc	agaaaatgga	taattagggg	180
ggccagcaaa	aatatcaagg	gtcaaataatc	gcacattttct	gtttaggcca	tctatggcct	240
tcatctcctc	tgaagtcaac	tgggaattcaa	acacctgcac	gttccgtctg	atgcgctgct	300
cattgtagct	cttgg					315

<211> 377

<213> Homo sapien

cctgccqtcg atgccaggga ggccgacagg accttctttt ccagcggggc cgatatattcc 60

```

aggggaacca ggaagacctc tgggtcccat gagaccaggc tccccagggc gaccagcatc 120
tccattaggt cctcggactc cagcagggcc acttgacca cgactaccag gagggcccat 180
gacgccagct ctgccatcag ctccaggaag accacgagaa ccaggactac ctctcagccc 240
aggaggtcct ggagggccgg cagatccagc tccccatta gggectctct ttccttcttc 300
accactggga ccaggaggac cttggggccc agcagagccg ggctcaccct tgttaccgct 360
ctctcctttg gagccag 377

```

<210> 1137

<211> 250

<212> DNA

<213> Homo sapien

<400> 1137

```

ctgttcaact tccaactcta aataggcacc attaaacaaa aaaccccagt attttaaatt 60
tctccagcac acattccagg atcaatgctc tgaactgtaa tcagctagta attcataacg 120
ggaatacagc cttagaatgg aagctatatt gcttccctgc cccctttctc ttacaattgg 180
agagtgtagg tattaagggg taaaaagtca gaggaagaat aattaaaaag aaaaatgccc 240
aaagctgcag 250

```

<210> 1138

<211> 511

<212> DNA

<213> Homo sapien

<220>

<221> misc_feature

<222> (1)...(511)

<223> n = A,T,C or G

<400> 1138

```

tcgaccaggt cctcctgggc catctggtcc ccgaggtcag cctggtgtca tgggcttccc 60
cggtcctaaa ggaaatgatg gtgctcctgg taagaatgga gaacgaggtg gccctggagg 120
acctggccct cagggtcctc ctggaaagaa tggtgaaact ggacctcagg gacccccagg 180
gctactggg cctggtggtg acaaaggaga cacaggaccc cctggtccac aaggattaca 240
aggcttgctt ggtacaggtg gtccctcagg agaaaatgga aaacctgggg aaccaggtcc 300
aaaggggtgat cccgggtgcac ctggagctcc aggaggcaag ggtgatgctg gtgcccctgg 360
tgaacgtgga cctcctggat tggcaggggc cccaggaactt agaggtggag ctgggtcccc 420
tgggtcccga ngaggaaagg gtgctgctgg tcctcctggg ccacctggtg ctgctggtac 480
tcctggtctg caaggaatgc ctggagaaag a 511

```

<210> 1139

<211> 505

<212> DNA

<213> Homo sapien

<400> 1139

```

ctgtggactc cagcatgttt ctgataatta tgcaagcaac aattctgtag cctcaagtaa 60
gaccacctgt gaacttgatc attatctggc ccaaatatga agataaacta taactttgga 120
gtttgtttcc tatttgattt cacattctgc ttctaaatc agttttctaa atttgtcctg 180
caattaggca ttggtcaggg gtgaatggct cttttcacag agagtagcca accagagacc 240
tttgctttga tatcatcaac tgcagagaat gctgttgatg ggaatgctgg aagcagaaac 300
tttgtcatcg gaaaaacttt tcttgatatg atgagactca acatcaggat ccacagctta 360
aagatgggaa ttcaggtatg aaagaaaaca ggcaaggagg cactgaggga gaaagacaca 420
gactttatcg ctctgtggct cattgttact ggaatattct aaaactcttg ttcacatgct 480

```

505

<213> Homo sapien

ctgtagcttc	tgtgggaatt	ccactgctcg	ggcgtcaggc	tcaggtagct	gctggccgcg	60
tacttgttgt	tgtctgttt	ggagggttt	gtggtctcca	ctccgcctt	gacggggctg	120
ccatctgcct	tccaggccac	tgtcacagct	ccggggtaga	agtcactgat	cagacacact	180
agtgtggcct	tgttggcttg	gagctctca	gaggaggcg	gaaacagagt	gacagtgggg	240
ttggccttgg	gctgac					256

<213> Homo sapien

ccagggcccc	attctgtctg	tgggactgtg	ggttctcagt	ggaattgttg	cctttcttgt	60
cgtaggagaaa	tttgtgagac	atgtgaaagg	aggacatggt	cacagtcatg	gacatggaca	120
cgctcacagt	catgcacgtg	gaagtcatgg	acatggaaga	caagagcgtt	ctaccaagga	180
gaagcagagc	tcagaggaag	aagaaaagga	aacaagaggg	gttcagaaga	ggcgaggagg	240
gagcacagta	cccaaagatg	ggccagtgag	acctcagaac	gctgaagaag	aaaaaagagg	300
cttagacctg	cgtgtgtcgg	ggtacctgaa	tctggctgct	gacttggcac	acaacttcac	360
tgatgggtctg	g					371

<213> Homo sapien

<223> n = A, T, C or G

cctccacac	tgtcaaatgt	caactccacc	agcactgaga	caatgagtag	atgagaatgt	60
agaaagaggg	aaggtggtag	gtaaaggagc	ggaaggaaga	ggtgggga	gagggaaggt	120
ggtaggtaaa	ggagcggaa	gaagaggtgg	ggaaagaggg	aaggagagaa	gggaaggagg	180
gaagagaaa	aaggaagaaa	aggaaagcat	ggcccggcta	gagacaaagc	cagaggtgat	240
caggtcagca	gcaggagagg	ctcagaagg	agcctctcgg	gaagtgcagg	cngccatgag	300
qqctcgtttc	ag					312

<213> Homo sapien

ccagacgtgg	tggctcacac	ctgcaatccc	agcaccttag	gaggccgagg	caggaggatc	60
cttgaggtca	ggagttcgag	accagcctcg	ccaacatggt	gaaaccccat	ttctactaaa	120

atacaaaaaa	ttagccaagt	gtggtggcat	atgcctgtaa	tcccaactac	tcagaaggcc	180
gaggcaggag	aattacttga	acgcaggaga	atcactgcag	cccaggaggc	agaggttgca	240
gtgagccgag	attgcaccac	tgcactccag	cctgggtgac	tgagcaagac	tccatctcag	300
taaataaata	aataaataaa	aagcgctgca	gtagctgtgg	cctcacccctg	aagtcagcgg	360
gcccagg						367

<210> 1144

<211> 159

<212> DNA

<213> Homo sapien

<400> 1144

cctggaggag	cggccgcaca	cacagccagg	cgctaggctc	cctgcgggac	ctcgggaagg	60
gggaagagcg	tcaacgattt	acggagggtc	cagccgctgg	gtcagattga	gacaaaccat	120
tgtgtggttg	ggttcgggtc	agcaggctgg	agagggttc			159

<210> 1145

<211> 450

<212> DNA

<213> Homo sapien

<400> 1145

ccatgggtgt	ctggagcacc	ctgaaactgt	atcaaagttg	tacatatttc	caaacatttt	60
taaaatgaaa	aggcactctc	gtgttctcct	cactctgtgc	actttgctgt	tgggtgtgaca	120
aggcatttaa	agatgtttct	ggcattttct	ttttatttgt	aagggtggtg	taactatggt	180
tattggctag	aaatcctgag	ttttcaactg	tatatatcta	tagtttgtaa	aaagaacaaa	240
acaaccgaga	caaacccttg	atgctccttg	ctcggcgttg	aggctgtggg	gaagatgcct	300
tttgggagag	gctgtagctc	agggcggtgca	ctgtgaggct	ggacctgttg	actctgcagg	360
gggcatccat	ttagcttcag	gttgtcttgt	ttctgtatat	agtgacatag	cattctgctg	420
ccatcttagc	tgtggacaaa	gggggggtcag				450

<210> 1146

<211> 324

<212> DNA

<213> Homo sapien

<400> 1146

ccatacaggg	ctgttgccca	ggccctagag	gtcattcctc	gtaccctgat	ccagaactgt	60
ggggccagca	ccatccgtct	acttacctcc	cttcggggcca	agcacaccca	ggagaactgt	120
gagacctggg	gtgtaaatgg	tgagacgggt	actttggtgg	acatgaagga	actgggcata	180
tgggagccat	tggctgtgaa	gctgcagact	tataagacag	cagtggagac	ggcagttctg	240
ctactgcgaa	ttgatgacat	cgtttcaggc	cacaaaaaga	aaggcgatga	ccagagccgg	300
caaggcgggg	ctcctgatgc	tgga				324

<210> 1147

<211> 191

<212> DNA

<213> Homo sapien

<400> 1147

ccacgaaaaat	caatgagaag	ccacaggtga	tcgcggacta	tgagagcgga	cggggccatac	60
ccaataacca	ggtgcttggc	aaaatcgagc	gggccattgg	cctcaagctc	cggggaaagg	120
acattggaaa	gcccacgag	aaggggccta	gggcgaaatg	aacacaaagc	ctcgaaatca	180
gtgtgctcca	g					191

<210> 1148
 <211> 344
 <212> DNA
 <213> Homo sapien

<400> 1148
 ctgtccaatg acaacaggac cctcactota ctcagtgtca caaggaatga tgtaggaccc 60
 tatgagtgtg gaatccagaa cgaattaagt gttgaccaca ggcacccagt catcctgaat 120
 gtccctctatg gccagacga ccccaccatt tccccctcat acacctatta ccgtccaggg 180
 gtgaacctca gcctctcctg ccatgcagcc tctaaccacac ctgcacagta ttcttggtg 240
 attgatggga acatccagca acacacacaa gagctcttta tctccaacat cactgagaag 300
 aacagcggac tctatacctg ccaggccaat aactcagcca gtgg 344

<210> 1149
 <211> 329
 <212> DNA
 <213> Homo sapien

<400> 1149
 ctgaccact cactgggcgg gggcacaggc tctggaatgg gcactctcct tatcagcaag 60
 atccgagaag aataccctga tcgcatcatg aataccttca gtgtggtgcc ttcacccaaa 120
 gtgtctgaca ccgtggtcga gccctacaat gccacccctct ccgtccatca gttggtagag 180
 aatactgatg agacctattg cattgacaac gaggccctct atgatattctg ctccgcact 240
 ctgaagctga ccacaccaac ctacggggat ctgaaccacc ttgtctcagc caccatgagt 300
 ggtgtcacca cctgcctcgg tttccctgg 329

<210> 1150
 <211> 406
 <212> DNA
 <213> Homo sapien

<400> 1150
 ccagttatatt gcaagtggta agagcctatt taccataaat aatactaaga accaactcaa 60
 gtcaaaccctt aatgccattg ttattgtgaa ttaggattaa gtagtaattt tcagaattca 120
 cattaacttg attttaaaat cagttttgtg agtcatttac cacaagctaa atgtgtacac 180
 tatgataaaaa acaaccattg tattcctgtt tttctaaaca gtactaattt ctaacactgt 240
 atatatcctt cgacatcaat gaactttgtt ttcttttact ccagtaataa agtaggcaca 300
 gatctgtcca caacaaactt gccctctcat gccttgctc tcaccatgct ctgctccagg 360
 tcagccccct tttggcctgt ttgttttgtc aaaaacctaa tctgct 406

<210> 1151
 <211> 346
 <212> DNA
 <213> Homo sapien

<400> 1151
 ctgcgtgagt accaggagct gatgaacgtc aagctggccc tggacatcga gatcgccacc 60
 tacaggaagc tgctggaggg cgaggagagc cggctggagt ctgggatgca gaacatgagt 120
 attcatatga agaccaccag cggctatgca ggtggtctga gctcggccta tgggggacctc 180
 acaagccccg gcctcagcta cagcctgggc tccagctttg gctctggcgc gggtccagc 240
 tccttcagcc gcaccagctc ctccagggcc gtggttgtga agaagatcga gacacgtgat 300
 gggaagctgg tgtctgagtc ctctgacgtc ctgcccgaagt gaacag 346

<210> 1152
 <211> 427
 <212> DNA
 <213> Homo sapien

<400> 1152
 ctggactgct gtacatcaag gacagattaa ctggaaaaca tatgttcctt atgcgtgac 60
 gagagccatt cagaaaagac ttcctttgtg ttcagcctat acttttccat atggtatacc 120
 ttgaaaaaaa ttagcacacc atgggttattt ttctacctt tataaaagac agagcctgtt 180
 tactcattta gaagatagag aaaattgggtc taaaattgaa catcctagat tcacactccc 240
 aagtcactta aggtgatttg atggtgagga aaatgattga cagagcccaa caatgatctc 300
 aggaattaca ttttccaaca gacaaaaaaa tgttttcatg tagcagcaat gcagatttgg 360
 tgaatattta atatataatt tagtatgtat ttcactttat gactgacaat taaaaaatat 420
 tgtttgg 427

<210> 1153
 <211> 331
 <212> DNA
 <213> Homo sapien

<400> 1153
 ctggccggcg gtgcagatct ggagtccagc ctcagggatg cgctactttc cattctctgc 60
 attgaacatt cgttctgtca gcatccgctc cagcttccact gcatcagcgg caaacttgcg 120
 gatcccgtca gagagcttct ccacagccat ctggctcctcg ttgtgcaacc aacggaaaga 180
 cttctcatcc aggtggattt tttccaggtc actggcttgg gctggggggac aagaaccagc 240
 cttccatgcc tgctccatgt ccctgccac cttggccct tgggctcagg gctgaaccg 300
 ctgcacccaa gcatctccca ccagggccag g 331

<210> 1154
 <211> 403
 <212> DNA
 <213> Homo sapien

<400> 1154
 ctgaactttc agatgaagtt gacttctact tgattgcagg attcagggtt tctcagatgt 60
 taatacagag tcaaaagcgg tggataaaac cttgcaaatt gcttgtgctt gttccaggct 120
 gttgactga taaaccaca ggctgtattc ctcattgctt gcatctgtgg tcttcagagc 180
 cagtaagctt tttcccgccc ccagaccgtc atcgtaacac accatccgga ttattaagta 240
 gagagcatgc ctgtgcaaaa catcatattg atctgatgtt gatactttta tgccatactt 300
 ggaaactccc ataataaatt cttcctccgg aggaacaaaa ggcaactttc catcttgctg 360
 ggcaacgtct atataattta tcagggtctaa tggcccttca agg 403

<210> 1155
 <211> 491
 <212> DNA
 <213> Homo sapien

<400> 1155
 cctccctctc agagcttgcc ccagggactc tctggccctc agggttcaat gtattctgac 60
 caaggccaag ctttcctggg gctcaggga aatcacactt tgctacccga agctgtatcc 120
 cctcagatgc caggaaggcc gtgatcatct gactccaccc tcctgagaca cattctctcc 180
 ctgactgtcc tgttctaagt cagcggagca ccttaggatg gaggggtgga ggcgaggcca 240
 gatgcagcct ctgtgaacag gtgcctggag gctgggaaat gaccctgaga ggcgaggaca 300
 cagcaaccgt gggcttaagg tgacctgag agcaagcttg gcccaacttta caattctgtt 360

```
<210> 1156
<211> 586
<212> DNA
<213> Homo sapien
```

```
<210> 1157
<211> 392
<212> DNA
<213> Homo sapien
```

```
<220>
<221> misc_feature
<222> (1)...(392)
<223> n = A,T,C or G
```

```
<210> 1158
<211> 375
<212> DNA
<213> Homo sapien
```

<400> 1158						
gggaaaaata	atattattcc	tcaaatgac	agcacattca	gaagcaggac	agaggagctc	60
tgatgacatc	tctgggggac	tcaaagcggc	cctcattttc	tggtattttc	ccaggtgatt	120
ctcttccaac	ctgtgagtc	tgtctctttt	cctcccatct	gaagtttgag	acatcctctg	180
ccacaaggaa	agccaccaat	accagcccaa	agagccacca	gagaggaacc	aaaccacatg	240
catcaagtta	taggaaggat	gcaagaaggg	aaattaggaa	ggaaagggag	gagtttagtt	300
ggcattctgg	ggcatgctaa	catgagggcg	atggtctctc	tccaagtcgc	tggacatatc	360
ccttttcttt	ccagg					375

```
<220>
<221> misc_feature
<222> (1)...(361)
<223> n = A,T,C or G
```

```
<210> 1160
<211> 142
<212> DNA
<213> Homo sapien
```

```
<210> 1161
<211> 193
<212> DNA
<213> Homo sapien
```

```
<210> 1162
<211> 265
<212> DNA
<213> Homo sapien
```

<210> 1163
<211> 337

<213> Homo sapien

<221> misc feature

<223> n = A, T, C or G

ctgcagagtg	ggganaggct	tttgccacta	gaaacttcca	ggatgcacga	gatcaaggaa	60
ttaagtctgt	aacaaaataa	caggatgctc	tgtgaagtcc	aaagaattgc	ttgaggcaaa	120
ctgcagagct	ccatgagatc	agcaacccca	agagctttta	caccgcgga	cacggtttaa	180
taggaaaaaa	atctcctata	ctgnntattc	anaaccaaat	gaanagaaat	gtcaaaggag	240
tcggaaacaa	tatgtcaaat	tangtaaat	cctgacctga	cccanatttt	gcngaacatt	300
tgtatcctaaa	ctgtctcttc	cacgtcctta	ggatcac			337

<211> 368

<213> Homo sapien

<221> misc feature

<223> n = A, T, C or G

cagacgtgg	tggctcacac	ctgcaatccc	agcaccttag	gaggccgagg	caggaggatc	60
cttgagggtca	ggagttcgag	accagcctcg	ccaacatggt	gaaaccccat	ttctactaaa	120
aatacaaaaa	attagccaag	tgtggtggca	tatgcctgta	atcccaacta	ctcagaaggc	180
cgaggcagga	gaattacttg	aacgcaggag	aatcactgca	nccangagg	canaggttgc	240
antgagccga	gattgcacca	ctgcactcca	gcctgggtga	cagagcaaga	ctccatctca	300
gtaaataaat	aaataaataa	aaagcgctgc	agtagctgtg	gcctcaccct	gaagtcagcg	360
ggcccagg						368

<211> 267

<213> Homo sapien

<221> misc feature

<223> n = A, T, C or G

ctgggaagga	ggctcctcgg	ccttctcctg	tttgtcatcc	tctcatcag	actcgacctc	60
catctcaact	tctcactct	ccccaaactt	ttcatagcgc	tctgaatga	ggattcgggc	120
cccagctcc	tctggcgtgg	tggggggagg	gaagtccct	tgctcattgg	gttggaagnc	180
cactgtttcc	accaccacaa	aatcatgcc	ntcnatctga	gcataggcca	cccgntcctt	240
ctcttctcc	nnttctctt	ttctctt				267

<211> 433

<212> DNA
<213> Homo sapien

<220>
<221> misc_feature
<222> (1)...(433)
<223> n = A,T,C or G

<400> 1166
ctgtctgtac actttttctt gggggaagag ttcttgtctt cagtttactg cagtaggggtt 60
cctggctctg ttacatgctc atgtgttccg gaagaacaca tgaaatatca tcccacggat 120
gacgatacag cccctgcttc ancctcttct gatcaagata gtgtccaatg aaccccatac 180
tccttcccag cacaaagatg ccattgaggg ctccaatgtc aatatattca tcagcttccct 240
ccctgcaaca cacatcaact tgtagtttta aaagggtcac gtgactgcc tccctcccac 300
agacagtact actactgccc aanaatgaga agaaaagggg tgctctgggt ggtngcatta 360
caggcaattt ttgttntctt nnttatacct ctccttattt tncaaanttt ctattatgag 420
tntgcattac ttt 433

<210> 1167
<211> 362
<212> DNA
<213> Homo sapien

<400> 1167
cctctggctc tttcttcagc caettotcca gctcctgcag gttctgggtc gagtagtcag 60
tgacgacgat ctccctaaag gattcacaaag cagagaggag ctgatagata gtggggccag 120
agccgatgtc aatcagcagg tctcccttca caccgtctag gcagaatata ttgaaaagat 180
ttttcagaag gtgcttaaga atctggcttt ctgcagagtg cctagaacca aacttgtaat 240
atTTTTctag gtaatcccga ggggtaaaat ggcttagata ggtgtccttg gaggtgaagc 300
ctgattccat tatgtctcac ttccgtacca ctggagcact gccctccttc tctttcctcc 360
ag 362

<210> 1168
<211> 459
<212> DNA
<213> Homo sapien

<220>
<221> misc_feature
<222> (1)...(459)
<223> n = A,T,C or G

<400> 1168
gcagtcattg ggcccaggac catgccactg gccctgctcc cccagccgca gcctcacctg 60
caggtgctcc tcgatgtcct tgcggtcgta ggtgatgcca ctgggcgtga tgcaaggctc 120
ccgcatcagc tcaaagctga tcttgccaca caggtagtcg gggatgtctc gcttctgtgg 180
cacaggggca cacggtcaga ggctgaaaag gggcactgca cgagcacctg ccagccatcg 240
gcagcaagcg acacacactc accttctct tctcatccac ctgagaaaaa agctcgtcca 300
tgtccgcat gtacttgctc tgtgaagagt tgagtgtgt gcttggggga gacacccac 360
ctccctcctn catggggcac anacccaaca caaggcgggg atgctnccac gccacgtgca 420
cacacacaga cccacatgtg ggtggggggc accctcacg 459

<210> 1169
<211> 386

<212> DNA

<213> Homo sapien

<400> 1169

ccaggccacc	tgtgcggggc	tcctcgatgt	ggaaggttcg	ggtgaggaga	ttgtagaagg	60
agccgtagca	cacggccacc	acagtgcacg	tgaggcagat	cacgctgtag	ggcatgctga	120
agtccggtgt	cggcagggttc	accagcagcg	gtcccggtgta	gagccgcaca	aagtagttag	180
agccatcaga	gactgggaac	aggctgttga	agaggggact	ctcttcccag	tccactggct	240
tggctgctac	catgctgggc	acaagggcgc	tgaggacaga	tgggctgaca	tagaagccat	300
ggttaggatc	tggcgtgtac	tcggtccact	tcagcagcgc	ccgctcaaac	tggatggaaa	360
ccttggtgac	tgagttggcc	ggccag				386

<210> 1170

<211> 480

<212> DNA

<213> Homo sapien

<400> 1170

ctattttctct	gttagtgttt	aaccaaccat	ctgttctaaa	agaagggctg	aactgatgga	60
aggaatgctg	ttagcctgag	actcaggaag	acaacttctg	cagggtcact	ccctggcttc	120
tggaggaaag	agaaggaggg	cagtgtctca	gtggtacaga	agtgagacat	aatggaatca	180
ggcttcacct	ccaaggacac	ctatctaagc	cattttaacc	ctcgggatta	cctagaaaaa	240
tattacaagt	ttggttctag	gcactctgca	gaaagccaga	ttcttaagca	ccttctgaaa	300
aatcttttca	agatattctg	cctagacggt	gtgaaggag	acctgctgat	tgacatcggc	360
tctggcccca	ctatctatca	gtcctctct	gcttgtgaat	cctttaagga	gatcgtcgtc	420
actgactact	caggaccaga	acctgcagga	gctggagaag	tggctgaaga	aagagccaga	480

<210> 1171

<211> 317

<212> DNA

<213> Homo sapien

<400> 1171

cctcagcagc	cctgccacgg	atctgcccga	ttctttcgca	tcaagaagtt	gatcttgcca	60
gccattttcca	tgttgtagat	ccgccggcac	ctttcatagc	tttccctctg	tcgccggcgg	120
catggcttct	cataataccg	ccgatgctta	atgtcctcaa	tgagcccatc	catagtgagg	180
attctgttta	gggtcctgta	tgcgctttcc	acgttccctt	cctgtaccat	cacagtctcg	240
gcgatgaact	tcagatgttt	tgccatgacc	ttggatttaa	accttcactc	tgtagagcct	300
cgcgcgctca	gtaccta					317

<210> 1172

<211> 202

<212> DNA

<213> Homo sapien

<220>

<221> misc_feature

<222> (1)...(202)

<223> n = A,T,C or G

<400> 1172

ggcaacggga	ggaacagcag	cagaggcagc	angagcagga	ggagcgtgaa	cgagaagagc	60
anccgcatn	ngctgcncctc	agtgaccgan	agaagagagc	tctggctgca	nagcgccgac	120
tcgctgcccc	gttgggagcc	cctacctctc	caatccctga	ctctgcaatc	gtcaatactc	180

gacgctgctg gagttgtggg gc

202

<210> 1173

<211> 173

<212> DNA

<213> Homo sapien

<400> 1173

ctgcctgggt	tgtggccgcc	ctagcatcct	gtatgccac	agctactgga	atccccgctg	60
ctgctccagg	ccaagcttct	ggttgattaa	tgagggcatg	gggtgggtccc	tcaagacctt	120
cccctacctt	ttgtggaacc	agtgatgcct	caaagacagt	gtcccctcca	cag	173

<210> 1174

<211> 301

<212> DNA

<213> Homo sapien

<400> 1174

ccaagagcta	caatgggcag	cgcacacagac	agaacgtgca	ggtttttgag	ttccagttga	60
ctgcggagga	catgaaagcc	atagatggcc	tagacagaaa	tctccactat	tttaacagtg	120
atagttttgc	tagccaccct	aattatccat	attcagatga	atattaacat	ggagagcttt	180
gcctgatgtc	taccagaagc	cctgtgtgtg	gatggtgacg	cagaggacgt	ctctatgccg	240
gtgactggac	atatcacctc	tacttaaata	cgctcctgtt	agcgacttca	gtcaactaca	300
g						301

<210> 1175

<211> 537

<212> DNA

<213> Homo sapien

<400> 1175

cctgcagggc	tggccgtag	gagaagggtca	gggcccaggg	cttcagcagg	gggcacttgt	60
taatggcatt	gaggttgatg	gacgcctcct	cctcactctg	gcctccagac	aggaagggtga	120
tcccagtgac	agcggggggc	actgtgcggc	gcagcgctgt	gacggtcggc	atggcaatct	180
cctcatgaga	aaacttctga	gtgcaagcat	ggcctggggg	gacctgttg	ggcttcagca	240
aggtgccttc	caggtagatg	tggtggtcac	tcagagcctt	gtagacagca	gccagcacct	300
tctcggtcac	atactggcag	cgcttcaagt	catgggtccc	atcaggagg	atctcaggct	360
ccacgatggg	cacaatgcc	ttctgctggc	agatactggc	ataacggggc	agaacattgg	420
cattttccat	gatggcgagg	gctgaggggg	tgtgttcccc	aatcttcagc	acacaacgcc	480
acttggcgaa	gtcagctccg	tccttcttgt	actgggcaca	gcgctcagac	agcccat	537

<210> 1176

<211> 384

<212> DNA

<213> Homo sapien

<220>

<221> misc_feature

<222> (1)...(384)

<223> n = A,T,C or G

<400> 1176

ctgacaaaaa	atgtgaaatt	tccacaaaat	atccaaactta	tgtgactaaa	cgcagtagtt	60
tttttaaaag	gggagataga	aaataaatgg	ttttgttgga	gtgcatttta	gtaagccttt	120

```
gcagtaaaat gacggttgta actactaaac caaatttagt tttcacagca tggttttgtt 180
gttttcccct tgtttttcag aggtaaaattt tgcattatat ccttcagtat tttaacacta 240
ttttggcagt ttacacatta ctttttgntt ttccttcctt tttgngaaat gtattaagtt 300
gtggttctta ttgaaacagt attatataat gttngcttaa ttatatcatg tgatgctcan 360
ntctattntg atttattcat tagt 384
```

<210> 1177

<211> 562

<212> DNA

<213> Homo sapien

<220>

<221> misc_feature

<222> (1)...(562)

<223> n = A,T,C or G

<400> 1177

```
ccaacaacat gcaggaagct cagagtatcg atgaaatcta caaatacgac aagaaacagc 60
agcaagaaat cctggcgggc aagccctggg ctaaggatca ccattacttt aagtactgca 120
aaatctcagc attggctctg ctgaagatgg tgatgcatgc cagatcgagg ggcaacttgg 180
aagtgatggg tctgatgcta ggaaagggtg atggtgaaac catgatcatt atggacagtt 240
ttgctttgcc tgtggagggc actgaaaccc gagtaaatgc tcaggctgct gcatatgaat 300
acatggctgc atacatagaa aatgcaaaac aggttggccg ccttgaaaat gcaatcgggt 360
ggtatcatag ccaccctggc tatggctgct ggctttctgg gattgatgtt agtactcaga 420
tgctcaatca gcagttccag gaaccatttg tagcagtggg gattgatcca acaagaacaa 480
tatccgcagg gnaaagtga tcttggcgcc tttaggacat acccaaaggg ctacaaacct 540
nctgatgaan gaccttctga gt 562
```

<210> 1178

<211> 353

<212> DNA

<213> Homo sapien

<220>

<221> misc_feature

<222> (1)...(353)

<223> n = A,T,C or G

<400> 1178

```
cgcgtctgga tggccgaatc attcgcacag actgggacgc aggctttaag gagggcaggc 60
aatacggccg tgggcatctt gggggccagg ttcgggatga gtatcggcag gactacnatg 120
ctgggagagg aggctatgga aaactggcac agaaccagtg agtggtgaga gctctgtcag 180
tgacaaacac tcctttggcc tgttgaattt gctgaagaac atcacctaaa gtctgcacac 240
gagcccatth ttaccaagat ttgatcagtg tctttactga gctggaagcc tctgaaagtt 300
attaaaggac agaatccaaa agaatgcctt taattcttgt ctgagaatct tgg 353
```

<210> 1179

<211> 288

<212> DNA

<213> Homo sapien

<400> 1179

```
ccaatgggat cctcaagggt cctgccatca atgtcaatga ctccgtcacc aagagcaagt 60
ttgacaacct ctatggctgc cgggagtccc tcatagatgg catcaagcgg gccacagatg 120
```

cgcgtctctg	acactgtgat	catgataggg	gttcaaacag	aaagtgcctg	ggccctcctt	60
ctaagtcttg	ttacaaaaaa	aaggaaaaag	aaaagatctt	ctcagttaca	aattctggga	120
agggagacta	tacctggctc	ttgccctaag	tgagaggtct	tccctcccg	acaaaaaat	180
agaaaggctt	tctatttcac	tggcccaggt	agggggaagg	agagtaactt	tgagtctgtg	240
ggcctcattt	cccaggtgcc	ttcaatgctc	atcaaaacca	ggcatgggga	aggccctggc	300
aaactgctcc	accggttgcc	tgaggttgg				329

<210> 1183
 <211> 198
 <212> DNA
 <213> Homo sapien

<400> 1183
 cctgacagac agaagggcctt ggagatTTTT tttctttaca attcagtctt cagcaacttg 60
 agagctttct tcatgttgct aagcaacaga gctgtatctg caggttcgta agcatagaga 120
 cgatttgaat atcttccagt gatatcggt ctaactgtca gagatgggtc aacaaacata 180
 atcctgggga catactgg 198

<210> 1184
 <211> 224
 <212> DNA
 <213> Homo sapien

<400> 1184
 ctggaggtgc ctcagaaggt gcattctgct tcttgcaggg gcttgaaaca ccaaggcact 60
 ccagggatcc tggagtcaaa gcagcagccc cgggtgttgct actccttggg ggtgacatgg 120
 gggtagccgc agtccaccct gtccttggct ggcacggcac actggtttgc agacaggccc 180
 acgtactcct cagcagagct ggaggacagc aaggccagga ccag 224

<210> 1185
 <211> 367
 <212> DNA
 <213> Homo sapien

<400> 1185
 cctttttacag atgtcagctt tctactggcct ccatgcacaa cctcccacta ccacccaatc 60
 tgcctgccac agcaaagtgc aggcaccctg ggccccctgg aggatgcggg caggggctac 120
 agggcatcca ggatgtggct gatcttgggt accagctcct ggcgctttcc tgagatgagc 180
 ttctcattct caatgtacgt gtctttcttg agcttgccag ccaccaggcg ctcagcctcc 240
 accgccgact tcagcaccag ctcttggacc tgtgcatcca gcttctgcat ttcgtcact 300
 ctgtgcgaca gatcagagcc ctctgtcttc agcctggact gcagcagtgc aatctcactg 360
 gtcaagg 367

<210> 1186
 <211> 188
 <212> DNA
 <213> Homo sapien

<400> 1186
 ccattaagcg gatgctggag atgggagcta tcaagaacct cacgtccttc cgacctgggc 60
 aagagctgta gcctgtcggg tgcctactct gctgtctggg tgacccccat gcgtggctgt 120
 ggggggtggct ggtgccagta tgaccactt ggactcacc cctcttgggg agggagtcct 180
 gggcctgg 188

<210> 1187
 <211> 379
 <212> DNA
 <213> Homo sapien

<400> 1187
 gttgatgcta ctctgaagtc tctcaacaac cagattgaga cccttcttac tcttgaaggc 60

```

tctagaaaaga gccagctcg cacatgccgt gacttgagac tcagccaccc agagtggagc 120
agtggttact actggattga ccctaaccaa ggatgcacta tggatgctat caaagtatac 180
tgtgatttct ctactggcga aacctgtatc cgggcccacac ctgaaaacat cccagccaag 240
aactggtata ggagctccaa ggacaagaaa cacgtctggc taggagaaac tatcaatgct 300
ggcagccagt ttgaatataa tgtagaagga gtgacttcca aggaaatggc tacccaactt 360
gccttcatgc gcctgctgg 379

```

<210> 1188

<211> 384

<212> DNA

<213> Homo sapien

<400> 1188

```

cgcgtcggac tgcagccagt ccgtttcctt tctttagcca gccatcctgg tactgtagtt 60
taggggttga tgggtggtga aattgatttc tggctgggta ctaagggtgcc tgctagccat 120
tgtataaaat taaaacatga agaataattt ttttttgagc atggctagtg gatttaaaac 180
aacacatacc tgtcactgct ggagtcaaac ttataaaaag ccttaagtgg aaagtgttcc 240
agacggagac tctgagttaa tagaggagta gaagctggtg tttaaagttcc cacgacgcac 300
atggctttgc cagaaactct gtttaatgat cgggcctttca cctcttcact tatccttagt 360
cccagtagcc aggatacctg atgg 384

```

<210> 1189

<211> 419

<212> DNA

<213> Homo sapien

<220>

<221> misc_feature

<222> (1)...(419)

<223> n = A,T,C or G

<400> 1189

```

ggaaaaacca gccactgctt tacaggacag ggggttgaag ctgagccccg cctcacaccc 60
acccccatgc actcaaagat tggattttac agctacttgc aattcaaaat tcagaagaat 120
aaaaaatggg aacatacaga actctaaaag atagacatca gaaattgttg agttaagctt 180
tttcaaaaaa tcagcaattc cccagcgtag tcaagggtgg aactgcacg ctctggcatg 240
atgggatggc gaccgggcaa gctttcttcc tcgagatgct ctgctgcttg agagctattg 300
ctttgttaag atataaaaag gggtttcttt ttgtctttct gtaaggtnna cttccagctt 360
ttgattgaaa gtccctagggt gattctatct ctgctgtgat ttatctgctg aaagctcag 419

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<210> 1190

<211> 173

<212> DNA

<213> Homo sapien

<400> 1190

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ccaggtactg gcacatcatg ctctggatgg ggggtggtgg gtccctgtagg cagagaaaca 60
ggaaattgtc gtagtcagta tcgagcagcg tggcctcgtt cggccacgta tagttgatct 120
tgaacttctt tggattctca gtcttctctc caaggacctt cttctcaaca cag 173

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<210> 1191

<211> 341

<212> DNA

<213> Homo sapien

<400> 1191
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 agttgcagca ctgagtgggc aaaatacatt tctggggccac ctcaggggaac ccatgcatct 120
 gcctggcatt taggcagcag agccctcgac cgtccccccac agggctctgc ctcacgtcct 180
 catctcattt ggctgtgtaa agaaatggga aaagggaaaaa ggagagagca attgaggcag 240
 ttgaccatat tcagttttat ttatttattt ttaatttggt cttttctcca agtccaccag 300
 tctctgaaat tagaacagta ggcggtatga gataatcagg a 341

<210> 1192

<211> 324

<212> DNA

<213> Homo sapien

<400> 1192
 ttggaggttg gcggcgcggg gctgaaggct agcaaaccga gcgatcatgt cgcacaaaaca 60
 aatttactat tcggacaaat acgacgacga ggagtttgag tatcgacatg tcatgctgcc 120
 caaggacata gccaaagtgg tccctaaaac ccatctgatg tctgaatctg aatggaggaa 180
 tcttggcggt cagcagagtc agggatgggt ccattatatg atccatgaac cagaacctca 240
 catcttgctg ttccggcgcc cactacccaa gaaaccaaag aaatgaagct ggcaagctac 300
 ttttcagcct caagctttac acag 324

<210> 1193

<211> 521

<212> DNA

<213> Homo sapien

<400> 1193
 ctgctttgtt ttctgttggc agtggaggga caagggtgaga ggagccaggg gtagtcatga 60
 acaccagtgg gttctgcctt gggcagctcc ccaccttctt taagagagta ctgtgtctca 120
 gctccagcag tctcaactgg gaagaccag gactcctgct cttttctcta atccctggga 180
 gacgaggtcc agctaaggta gagtaagcag tcagtgacca ggcaggctgg tttgggaggt 240
 cactgcctgg aggacgggat cttgtattct tcggaagatg gctgggaaat tcttccctcc 300
 attacgtaga actttcttcc cctcctcagt tgagggtgcct agatgtccca caacgggggtc 360
 ttcactcagg tcctccagag gcacacgctc aaacagtggg tgctcttcga aatgagtga 420
 catccagtgc tgtagctcca gcacatcggg tatggtatac accagcccct gcataggcaa 480
 aatcacccta gacaggaggc tgcattgcaac gtcagcagcc a 521

<210> 1194

<211> 208

<212> DNA

<213> Homo sapien

<400> 1194
 ccagtgacta gaaggcgagg cgccgcggga ccatggcggc ggcggcggac gagcggagtc 60
 cagaggacgg agaagacgag ggagaggagg agcagttggg tctggtggaa ttatcaggaa 120
 ttattgattc agacttcctc tcaaaatgtg aaaataaatg caagggtttg ggcattgaca 180
 ctgagaggcc cattctgcaa gtggacag 208

<210> 1195

<211> 499

<212> DNA

<213> Homo sapien

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<210> 1196
<211> 455
<212> DNA
<213> Homo sapien
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<210> 1197
<211> 444
<212> DNA
<213> Homo sapien
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<210> 1198
<211> 450
<212> DNA
<213> Homo sapien
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<400> 1198						
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aggcatttaa	agatgtttct	ggcattttct	ttttatttgt	aagggtggtg	taactatggt	180
tattggctag	aaatcctgag	ttttcaactg	tatatatcta	tagtttgtaa	aaagaacaaa	240
acaaccgaga	caaacccttg	atgctccttg	ctcggcgttg	aggctgtggg	gaagatgcct	300
tttgggagag	gctgtagctc	agggcgtgca	ctgtgaggct	ggacctgttg	actccgcagg	360
qqqcatccat	ttagcttcag	gttgtcttgt	ttctgtatat	agtgacatag	cattctgctg	420

ccatcttagc tgtggacaaa ggggggtcag

450

<210> 1199

<211> 294

<212> DNA

<213> Homo sapien

<400> 1199

agtcacagtt	gcacctattc	aaaactagct	ttaaagtgag	ctatTTTTTaa	acttcataaa	60
aatattcatg	atTTTattag	TTTgaatatt	tctacaagat	tcgggtgggc	TTTTcTTTa	120
ggTgaaaaca	gctatccact	cctgtggcct	tataactcag	gaaatgctgg	ggatgcaaac	180
gtgcaaaagg	cagggggaag	ctgcccaggc	tgagactgga	gcagctagga	gtgtgcttgg	240
ggaacgggag	ctgagatccc	ggagcagaaa	tggtcagccg	tgctctggag	cagg	294

<210> 1200

<211> 258

<212> DNA

<213> Homo sapien

<400> 1200

agctacctaa	gaacagctaa	aagagcacac	ccgtctatgt	agcaaaatag	tgggaagatt	60
tataggtaga	ggcgacaaac	ctaccgagcc	tggtgatagc	tggttgcca	agatagaatc	120
ttagttcaac	TTTaaTTTg	cccacagaac	cctctaaatc	cccttgtaaa	TTTaaCTgtt	180
agtccaaaga	ggaacagctc	TTTggacact	aggaaaaaac	cttgtagaga	gagtaaaaaa	240
TTTaaCacc	atagtagg					258

<210> 1201

<211> 403

<212> DNA

<213> Homo sapien

<400> 1201

ctgagctgct	gtctgctttg	gaaaaccggt	cctgccgctg	ccgatggatg	gaaatgcaat	60
ggatttcagc	ttcttatcat	cagccagggc	caagcagttt	ttcactgtct	ttccagaag	120
ttcttcacac	ttgtctgcac	cccaaactgg	actattacag	tgatcaca	acttggcagg	180
caggccatgg	cctgcgctga	cagcagctcc	agctacttcc	aagggcccg	TTTTTTccg	240
gagttccagg	acagcttcca	caaactcctt	gccacctttc	ttctccagcg	tgTTTcctag	300
gtcatcttta	aggtcaatgt	cagcattgg	aggattgatt	atggcctcca	cctcaaagcc	360
ggctaaatta	ctgatttcac	tgtgaataag	gttcggcttc	tgg		403

<210> 1202

<211> 325

<212> DNA

<213> Homo sapien

<400> 1202

ctgaacctgc	gggagtcggc	caccatcacg	tgcctggtga	cgggcttctc	tccgcggac	60
gtcttcgtgc	agtggatgca	gagggggcag	cccttgctcc	cggagaagta	tgtgaccagc	120
gccccaatgc	ctgagcccca	ggccccaggc	cggtaacttcg	cccacagcat	cctgaccgtg	180
tccgaagagg	aatggaacac	gggggagacc	tacacctgcg	tggtggccct	tgaggccctg	240
cccaacagg	tcaccgagag	gaccgtggac	aagtcaccgc	gtaaaccac	cctgtacaac	300
gtgtccctgg	tcatgtccga	cacag				325

<210> 1203

<211> 518
 <212> DNA
 <213> Homo sapien

<400> 1203
 ctcaaccaca gtctgacacc agagcccact tccatcctct ctggtgtgag gcacagcgag 60
 ggcagcatct ggaggagctc tgcagcctcc acacctacca cgacctccca gggctgggct 120
 caggaaaaac cagccactgc ttacaggac aggggggtga agctgagccc cgcctcacac 180
 ccacccccat gcaactcaaag attggatttt acagctactt gcaattcaaa attcagaaga 240
 ataaaaaatg ggaacataca gaactctaaa agatagacat cagaaattgt taagttaagc 300
 tttttcaaaa aaccagcaat tccccagcgt agtcaagggg ggacactgca cgctctggca 360
 tgatgggatg gcgaccgggc aagctttctt cctcgagatg ctctgctgct tgagagctat 420
 tgctttgtta agatataaaa aggggtttct tttgtcttt ctgtaagggt gacttccagc 480
 ttttgattga aagtcctagg gtgattctat ttctgctg 518

<210> 1204
 <211> 352
 <212> DNA
 <213> Homo sapien

<400> 1204
 ggggaaagga ggtctcactg agcaccgtcc cagcatccgg acaccacagc ggccttctgc 60
 tccacgcaga aaaccacact tctcaaactc tcaactcaaca ctctctccc caaagccaga 120
 agatgcacaa ggaggaacat gaggtggctg tgctgggggc acccccagc accatccttc 180
 caaggtccac cgtgatcaac atccacagcg agacctccgt gcccgaccat gtcgtctggt 240
 ccctgttcaa caccctcttc ttgaactggt gctgtctggg cttcatagca ttgcctact 300
 ccgtgaagtc tagggacagg aagatggttg gcgacgtgac cggggcccag ga 352

<210> 1205
 <211> 250
 <212> DNA
 <213> Homo sapien

<400> 1205
 ctgttcaact tccaactcta aataggcacc attaaacaaa aaacccagc attttaaatt 60
 tctccagcac acattccagg atcaatgctc tgaactgtaa tcagctagta attcataacg 120
 ggaatacagc cttagaatgg aagctatatt gcttccctgc cccctttctc ttacaattgg 180
 agagtgtagg tattaaggga taaaaagtca gaggaagaat aattaaaaag aaaaatgccc 240
 aaagctgcag 250

<210> 1206
 <211> 275
 <212> DNA
 <213> Homo sapien

<220>
 <221> misc_feature
 <222> (1)...(275)
 <223> n = A,T,C or G

<400> 1206
 ctgctctcgn ngnctcactg gatggaccag cacttccgca cgacgcccct ggagaagaac 60
 gccccgtct tgctggccct gctgggtatc tggatcatca actgctttgg gtgtgagaca 120
 cagcccatgc tgccctatga ccagtacctg caccgctttg ctgcgtactt ccagcagggc 180

gacatggagt ccaatgggaa atacatcacc aaatctggaa cccgtgtgga ccaccnnaca 240
ggccccattg tgtgggggga gccagggacc aatgg 275

<210> 1207
<211> 182
<212> DNA
<213> Homo sapien

<400> 1207
ccatctcctg ctgcaagtcc agggcgacgt agcacagctt ctctttgatg tcgcgcacga 60
tttcccgctc ggccgtggtg gtgaagctgt agcctcgctc agtgaggatc ttcagaggt 120
agtcggtcag gtcccgcca gccaggtcca gacgcaggat ggcgtggggg agggcgtagc 180
cc 182

<210> 1208
<211> 260
<212> DNA
<213> Homo sapien

<220>
<221> misc_feature
<222> (1)...(260)
<223> n = A,T,C or G

<400> 1208
gctggttatg aactcctgac ctcaagtgat ctgccctcct cagcctccca aagtgtctggg 60
attataggca tgagccactg gaatttttct tttttttttt ctttcttttt tttttttttt 120
ttaaattgan acaaggtctg gctctatcgc ccangctgga gtgcagnggc accatntcgg 180
ctcactgcaa cctctgcctg ctgggctcga gccatcctcc cacctcagcc tcccaagtan 240
ttgggactag aggtatgcac 260

<210> 1209
<211> 487
<212> DNA
<213> Homo sapien

<400> 1209
aaaccactc caccttacta ccagacaacc ttagccaaac catttaccca aataaagtat 60
aggcgataga aattgaaacc tggcgcaata gatatagtag cgcaagggaa agatgaaaaa 120
ctataaccaa gcataatata gcaaggacta atccctatac cttctgcata atgaattaac 180
tagaaataac tttgcaagga gagccaaagc taagaccccc gaaaccagac gagctaccta 240
agaacagcta aaagagcaca cccgtctatg tagcaaaata gtgggaagat ttataggtag 300
aggcgacaaa cctaccgagc ctgggtgatg ctgggtgtcc aagatagaat cttagttcaa 360
ctttaaattt gccacagaa ccctctaaat ccccttgtaa atttaactgt tagtccaaag 420
aggaacagct ctttggacac taggaaaaaa ccttgtagag agagtaaaaa atttaacacc 480
catagta 487

<210> 1210
<211> 216
<212> DNA
<213> Homo sapien

<400> 1210
ccactcagct cagcgggcca cgtgccccta caagttggca gaagtggctg ccactgctgg 60

gtttgtgtaa	gagaggtgc	tgccaccatt	acctgcagaa	accttctcat	aggggctacg	120
atcggtagctg	ctagggggca	catagcgccc	atggatgtgg	taggtggggg	actcgctcat	180
aggatggtag	gtatcccggg	ctggaaagat	gtccag			216

<210> 1211

<211> 443

<212> DNA

<213> Homo sapien

<400> 1211

ccaaggtcag	aggctgatgc	aacaggccct	cttctcccca	gggccaggct	cctgtccagc	60
ctgggcactg	cccagagtga	tggcattggg	ccggatgctg	ttctgtctct	gcttggacac	120
cttcgcaaag	atttctttca	ggacagtctc	aaaggctagc	tcaacattgg	tagagtccag	180
ggctgaggtc	tccaggaaga	gcagtcatt	gttttcagcg	aacattcggg	cctcctcagt	240
gggcacttcc	cgggcctggc	tgaggctcact	tttgttaccc	acgagcatga	cgacgatcgt	300
ggcttcagca	tggcataga	gctccttcag	ccatcgctcc	accacagcat	aggtctgggtg	360
cttggttagg	tcaaacacca	ggagggcccc	cactgcacca	cgatagtacg	ccgaggtgat	420
ggctcggtac	cgctccaggc	cag				443

<210> 1212

<211> 526

<212> DNA

<213> Homo sapien

<400> 1212

actgaaaccc	gagtaaagtc	tcaggctgct	gcatatgaat	acatggctgc	atacatagaa	60
aatgcgaaac	agggtggccg	ccttgaaaat	gcaatcgggt	ggtatcatag	ccaccctggc	120
tatggctgct	ggctttctgg	gattgatgtt	agtactcaga	tgttcaatca	gcagttccag	180
gaaccatttg	tagcagtggg	gattgatcca	acaagaacaa	tatccgcagg	gaaagtgaat	240
cttggcgctt	ttaggacata	cccaaagggc	tacaaacctc	ctgatgaagg	accttctgag	300
taccagacta	ttccacttaa	taaaatagaa	gattttgggtg	tacactgcaa	acaatattat	360
gccttagaag	tctcatattt	caaactctct	ttggatcgca	aattgcttga	gctgttgggtg	420
aataaatact	gggtgaatac	gttgagttct	tctagcttgc	ttactaatgc	agactatacc	480
actggtcagg	tctttgattt	gtctgaaaag	ttagagcagt	cagaag		526

<210> 1213

<211> 359

<212> DNA

<213> Homo sapien

<220>

<221> misc_feature

<222> (1)...(359)

<223> n = A,T,C or G

<400> 1213

ccagccattg	cctgncattt	ggtagtatag	tatgattctc	accattattt	gtcatggagg	60
cagacataca	ccagaaatgg	gggagaaaca	gtacatatct	ttctgtcttt	agttttattgt	120
gtgctgggtc	aagcaagctg	agatcatttg	caatggaaaa	cacgtaactt	gtttaaaagt	180
ttttctggta	gcttttagctt	tatgctaaaa	aaaataatga	cattgggtat	ctatttcttt	240
ctaagactac	attantanga	aaataagtct	tttcatgctt	atgatttagc	tgttttgtgg	300
taattgcttt	ttaaaggaag	nnattaatat	cataagttat	tattaatatt	gtgaacnca	359

<210> 1214

<211> 428
 <212> DNA
 <213> Homo sapien

<400> 1214
 ccaagcttga ggcagcccta ggtgaggcca agaagcaact tcaggatgag atgctgcggc 60
 ggggtggatgc tgagaacagg ctgcagacca tgaaggagga actggacttc cagaagaaca 120
 tctacagtga ggagctgcgt gagaccaagc gccgtcatga gacccgactg gtggagattg 180
 acaatgggaa gcagcgtgag tttgagagcc ggctggcggg tgcgctgcag gaactgcggg 240
 cccagcatga ggaccagggt gagcagtata agaaggagct ggagaagact tattctgcca 300
 agctggacaa tgccaggcag tctgctgaga ggaacagcaa cctgggtggg gctgcccacg 360
 aggagctgca gcagtcgcgc atccgcacgc acagcctctc tgcccagctc agccagctcc 420
 agaagcag 428

<210> 1215
 <211> 414
 <212> DNA
 <213> Homo sapien

<400> 1215
 ctgaagcact cttcagagac tacgtccaca gacactgatg ctgaggcctt tcttgtaagt 60
 gaagaaaaag gaatgcagca aagaagagtt cgacattgga gtccttagtt ccatcaggat 120
 cccatttcgca gccttttagca tcatgtagaa gcaaactgca cctatggctg agatagggtgc 180
 aatgacctac aagattttgt gttttctagc tgtccaggaa aagccatctt cagtcttgct 240
 gacagtcaaa gagcaagtga aaccattttcc agcctaaact acataaaagc agccgaacca 300
 atgattaaag acctctaagg ctccataatc atcattaaat atgcccacaa tcattgtgac 360
 tttttatttt atatacagga ttaaaatcaa cattaaatca tcttatttac atgg 414

<210> 1216
 <211> 162
 <212> DNA
 <213> Homo sapien

<220>
 <221> misc_feature
 <222> (1)...(162)
 <223> n = A,T,C or G

<400> 1216
 cctggccgca ggggtccccg gtattgctgt tgctacgagg ttgggggggca gcgattgtcc 60
 tgtgggagcc accgttctcc tgggtcgggg accctcactt cttctggggg gtgctcannt 120
 tctgcatgcc ccgatcttg tccagcangc cagaaatgaa gg 162

<210> 1217
 <211> 392
 <212> DNA
 <213> Homo sapien

<220>
 <221> misc_feature
 <222> (1)...(392)
 <223> n = A,T,C or G

<400> 1217

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ctgaagtaga ggctggaact gaagctgaga ctgaggctga ggctgaaact ggagctaagg      60
gtgaggctgg aactggagct gaggttgagg ccagaactgg agctaaagtt gaggctggaa      120
ccggagctga ggttgaggct ggaactggag ttaaggttgc tggaaagtga gctgaggttg      180
aggctggaac tgaagctgag gttgaagggt gaagtggagc cgaagctaga ggtggaactg      240
aggctgaaga ctgtgcttgc tggatccctg tagcctgttt tttggcaaact cttggaggaa      300
gcttanaagt ctggcttctt cctttttcat ttgcattctt tttgttccag accttaaaaa      360
attaacgggg accatttttg tcaataatgc ag                                     392

```

<210> 1218

<211> 526

<212> DNA

<213> Homo sapien

<220>

<221> misc_feature

<222> (1)...(526)

<223> n = A,T,C or G

<400> 1218

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ctgagctttc agcagataaa tcacagcaga aatagaatca ccctaggact ttcaatcaaa      60
agctggaagt ccaccttaca gaaagacaaa aagaaacccc tttttatata ttaacaaagc      120
aatagctctc aagcagcaga gcatctcgag gaagaaagct tgcccggtcg ccatcccatc      180
atgccagagc gtgcagtgtc cacccttgac tacgctgggg aattgctgat tttttgaaaa      240
agcttaactt aacaatttct gatgtctatc ctttagagtt ctgtatgttc ccatttttta      300
ttcttctgaa ttttgaattg caagtagctg taaaatccaa tctttgagtg catgggggtg      360
gggtgtgaggc ggggctcanc ttcaaccccc tgtcctgtaa agcagtggct ggtttttctt      420
gagcccagcc ctgggaggtc gtggtangtg tggaggctgc agagctcctn cagatgctgc      480
cctcgctgtg cctcacacca nagaggatgg aagtgggctc tgggtgt                                     526

```

<210> 1219

<211> 382

<212> DNA

<213> Homo sapien

<400> 1219

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ctggccggcg gtgcagatct ggagtccagc ctcagggatg cgctactttc cattctctgc      60
attgaacatt cgttctgtca gcatccgctc cagcttccact gcatcagcgg caaacttgcg      120
gatcccgctc gagagcttct ccacagccat ctggctcctcg ttgtgcaacc aacggaaaga      180
cttctcatcc aggtggattt ttccaggctc actggccttg gccgccttg ctgagagcac      240
aggcaccagc ttggcgttgt cctgcagcag ctctcccagg agcttggttg agatggtgag      300
gaagtcacag ccggccagtg ctttgatctc gcccggtgtg cggaaggagg cgcccatgac      360
aatggttttg tagctaaact tc                                     382

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<210> 1220

<211> 127

<212> DNA

<213> Homo sapien

<400> 1220

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tcgacctcct tgaagcagac caagtatagc aagcctctaa aaggactact gagaaacaga      60
atcagaaact ctagaactct agttagggcc cttcagcagg gctgcagagc ctccctggat      120
accaggg                                           127

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<210> 1221

<211> 304
 <212> DNA
 <213> Homo sapien

<400> 1221
 ccaccccgga gatgacacga ggctcacatg actctagaca cttggtggaa agtgaggcga 60
 gaaaaacaat gacttgggcc aattacacga ctgcaaagct agagctgcca acagggctcc 120
 agggagcttg gcttctgtag aagttctaag gaagcggtag gaactccacg gcggtggggc 180
 gctaactagc agggacccct gcaagtgttg gtcggggggc tcgggctgcc tgagctgaca 240
 cgaggggagg ggtctgtgta gccaacaggt gaccgaaggg cttgcctgcc cacagcttac 300
 ttgg 304

<210> 1222
 <211> 309
 <212> DNA
 <213> Homo sapien

<400> 1222
 ctgtcgact cgtagctgca actcaactcaa cttgtcttta gcagcaattt ctgcatagtc 60
 attggcatgt tcacctacct ggatgtccgg gtgaactctc agcatgcctc cagcaaagag 120
 ggagaacttg gtggaattgg agtgaagaca gatctggtgc tcaccagggg tatgggaagt 180
 gaaagtgaac ctgccctcgg agccatactg ccggggccagg atgaccttgt cctctgggtc 240
 ctccacctcc acaaacatgc caagccccgg ggtggccggc tgggtactcct cccgctgctt 300
 gtcatacag 309

<210> 1223
 <211> 390
 <212> DNA
 <213> Homo sapien

<400> 1223
 cctggcctgg gagccctgtg cctactagaa gcacattaga ttatccattc actgacagaa 60
 caggtctttt ttgggtcctt cttctccacc acgatatact tgcagtcctc cttcttgaag 120
 attcttttggc agttgtcttt gtcataaccc acaggtgtag aaacaagggt gcaacatgaa 180
 atctctgttt cgtagcaagt gcatgtctca cagttgtcag tctgccactc cgagtttatt 240
 ggtgtttgtt tcctttgaga tccatgcatt tcttggttga atctcctgga actccctcat 300
 taggtatgaa atagcatgat gcattgcata aagtcacgaa ggtggcaaag atcacaacgc 360
 tgcccaggag aacattcatt gtgataagca 390

<210> 1224
 <211> 407
 <212> DNA
 <213> Homo sapien

<400> 1224
 ccttatgact acaacggccc acgagaaaaa tatggaatcg ttgattacat gatcgagcag 60
 tccgggcctc cctccaagga gattctgacc ctgaagcagg tccaggagtt cctgaaggat 120
 ggagacgatg tcatcatcat cggggtcttt aagggggaga gtgaccacgc ctaccagcaa 180
 taccaggatg ccgctaacaa cctgagagaa gattacaaat ttaccacac tttcagcaca 240
 gaaatagcaa agttcttgaa agtctccag gggcagttgg ttgtaatgca gcctgagaaa 300
 ttccagtcca agtatgagcc ccggagccac atgatggacg tccagggctc caccaggagc 360
 tcggccatca aggacttcgt gctgaagtac gcctgcccc ttggttgg 407

<210> 1225

<211> 250
 <212> DNA
 <213> Homo sapien

```
<400> 1225
ctgcagcttt gggcattttt ctttttaatt attcttcctc tgactttgta tcccttaata      60
cctacactct ccaattgtaa gagaaagggg gcaggggaagc aatatagctt ccattctaag      120
gctgtattcc cgttatgaat tactagctga ttacagttca gagcattgat cctggaatgt      180
gtgctggaga aatttaaaat actgggggtt tttgtttaat ggtgcctggt tagagttgga      240
agttgaacag                                     250
```

<210> 1226
 <211> 444
 <212> DNA
 <213> Homo sapien

<220>
 <221> misc_feature
 <222> (1)...(444)
 <223> n = A,T,C or G

```
<400> 1226
ccttttagct gttgctctgg gcaggggggtg ggggtgcggg ggcttacagt gggggccctt      60
agttggcaca ggttcggaag ggcccagggc agacatgaat tctcctgaga cttgaggtag      120
gttgcttcag ccagcccggg cggagaagaa gggcagagag cgaacatagg agtccagtcg      180
ggagcgaaag agctcacttt gcacagtttg gccagcggg cacaggggat tcttcaccac      240
cagctccaca tacagcgcac tgtagatgtg gtgcagcaca tctcggatgg gtcccacgcc      300
caagtcagta ttcataacaa ctttgatccc agtgggcgtc tcgtagtaat ggagtttgta      360
acggctagtt tggaaggcca ggaagccatc cttcatgtct agcggggaca tcttgctgac      420
aaacgancgg atagagaaga gcat                                     444
```

<210> 1227
 <211> 491
 <212> DNA
 <213> Homo sapien

```
<400> 1227
gttagcctta catgttgtgt agacttactt taagtttgca cccttgaaat gtgtcatatc      60
aatttctgga ttcataatag caagattagc aaaggataaa tgccgaaggt cacttcattc      120
tggaacacagt tggatcaata ctgattaagt agaaaatcca agctttgctt gagaactttt      180
gtaacgtgga gagtaaaaag tatcggtttt attctttgct gatgtccttt ctgcttgaaa      240
taacagtcac catacagcta aaggagagga gtttctttcc ttctaagtag gcagaaatgg      300
tatcattatg ttgccgtctt ccaatctccc agagctcgtt ctctagagaa tcaccttctt      360
tcgttttttt tttttttttg aggtagagtc tcactatggt gccagacta gccttgaact      420
cctgggctca agtgattctc cctcctcagc ctcccagagta gctggaacga actatagttg      480
caccactgca g                                     491
```

<210> 1228
 <211> 279
 <212> DNA
 <213> Homo sapien

```
<400> 1228
ctgggcggat ctgatcaact aggcaacatc atgtccggat atgagttcat caacaagttg      60
```

ctgcaactttt	ttttttttgc	aattacagag	tggtattcag	ttaacagaac	aacaattatt	60
tcgataaagc	tgcatacagag	acaactgaag	atgaaaaaac	taccatcccc	atatataact	120
aatttggtgct	gtgcaccaac	aagaacctgc	tttaaatttc	catgcccaatt	tacaaccccc	180
atactgtacc	aggcaagggt	agtggctatt	gaaaatacca	ccaggacagg	gctatctaaa	240

```
<210> 1233
<211> 312
<212> DNA
<213> Homo sapien
```

<400>	1233						
ctgagcgtac	ggccgcgttc	atcccagccg	cgggtgcccc	cacgttgatg	acagctacgt		60
tgcaattggt	ctttgggate	tgatcatccg	gcagcttgat	ggcaagtcgc	ttgtaggtgt		120
tcaggttgcc	cgcaaagctc	ctccctcgga	gtcgaaccgn	atnttgaaat	ctcctctcgt		180
ccatcgccct	ctgcacatcc	tgagtcatct	gcacgcactc	catcagcggc	aggcgcacgg		240
ngtggttccc	gttcagtgac	acgacgcaag	ctggggtgtc	cggggtggcc	tctagcaagg		300
cnatgactgc	ct						312

```
<400> 1234
ccggccgcgagg gcataaaagg cgccaggtga gggcctcgcc gctcctcccg cgaatcgag 60
cttctgagac caggggtgct cgcgcgtgc tcgcctcgc catgacttcc tacagctatc 120
gccagtcgtc ggccacgtcg tccttcggag g                                     151
```

```
<220>  
<221> misc_feature  
<222> (1)...(250)  
<223> n = A,T,C or G
```

```
<210> 1236
<211> 154
<212> DNA
<213> Homo sapien
```

<400> 1236

ctgatcctca	ctattgtggg	caccatcgct	ggcatcgta	ttctcagcat	gataattgca	60
ttgattgtca	cagcaagatc	aaataacaaa	acgaagcata	ttgaagaaga	gaacttgatt	120
gacgaagact	ttcaaaatct	aaaactgcgg	tcga			154

<210> 1237

<211> 375

<212> DNA

<213> Homo sapien

<400> 1237

ccactggatc	tttgggatta	aagctctgtt	ggatttgtac	ctcagaggaa	gatcaagtgg	60
ctgatccttt	ggactctgta	aagagcattc	ttctagtcag	aggggtggaat	ggcagcagca	120
actggaagaa	aatgagtttt	ttggtgcccc	cacccaagag	cacacacatg	ctgcactgtc	180
tcggaagca	gggccagcta	gagccaccat	gttcttcctt	acctcagttt	acctgcggcc	240
tgcgctgcac	tgcagatgcc	caccctgccc	tgggtctggc	cggcggaagc	tctgtccaag	300
gtccacacac	ctccaggttt	acgccaacat	ccttgtgccc	tccccacctt	ctcttccaac	360
gcattaggtg	cattg					375

<210> 1238

<211> 454

<212> DNA

<213> Homo sapien

<400> 1238

gtcaagatca	agttcaatat	catcgctctt	ctctatgact	acaaccccaa	cctggcaacc	60
tacatgaagc	cagagatgtg	ggggaagtgc	ctggactgca	tcaatgagct	gatggatata	120
ctgtttgcaa	atcccaacat	ttttgttgga	gagaatattc	cgggaagagag	tgagaacctg	180
cacaacgctg	accagccact	gcgtgtccgt	ggctgcattc	taactctggt	ggaacgaatg	240
gatgaagaat	ttaccaaaat	aatgcaaaat	actgaccctc	actccaagag	tacgtggagc	300
acttgaagga	tgaggcccag	gtgtgtgcca	tcatcgagcg	tgtgcagcgc	tacctggagg	360
agaagggcac	taccgaggag	gtctgccgca	tctacctgct	gcgcattcctg	cacacctact	420
acaagtttga	ttacaaggcc	catcagcgac	agac			454

<210> 1239

<211> 483

<212> DNA

<213> Homo sapien

<400> 1239

ctgccaggct	gaaaagaagc	ctcagctccc	acaccgccct	cctcaccgcc	cttcctcggg	60
agtcacttcc	actgggtggac	cacgggcccc	cagccctgtg	tcggccttgt	ctgtctcagc	120
tcaaccacag	tctgacacca	gagcccaatt	ccatcctctc	tgggtgtgagg	cacagcgagg	180
gcagcatctg	gaggagctct	gcagcctcca	cacctaccac	gacctcccag	ggctgggctc	240
aggaaaaacc	agccactgct	ttacaggaca	gggggttgaa	gctgagcccc	gcctcacacc	300
cacccccatg	cactcaaaga	ttggatttta	cagctacttg	caattcaaaa	ttcagaagaa	360
taaaaaatgg	gaacatacag	aactctaaaa	gatagacatc	agaaattgtt	aagttaagct	420
ttttcaaaaa	atcagcaatt	ccccagcgta	gtcaagggtg	gacactgcac	gctctggcat	480
gat						483

<210> 1240

<211> 358

<212> DNA

<213> Homo sapien

```
<210> 1241
<211> 194
<212> DNA
<213> Homo sapien
```

```
<210> 1242
<211> 316
<212> DNA
<213> Homo sapien
```

```
<210> 1243
<211> 275
<212> DNA
<213> Homo sapien
```

```
<210> 1244
<211> 235
<212> DNA
<213> Homo sapien
```

<400>	1244								
ctgctgcgct	tggataacaa	gtaattcaac	gcacgcactt	aacagaaatg	ttaaactata				60
acaagcacca	tttgaggatt	aacaggaaca	tttttttgaa	gatttcaaac	gaactcgact				120
ttcagataaa	tgtacctaa	agtatttata	aacagctcat	cggagcctct	atttgtcata				180
qacttttgag	ttgattgttg	ggaccacata	ataggacct	tttttttttg	tcttt				235

<210> 1248

<211> 640
 <212> DNA
 <213> Homo sapien

<220>
 <221> misc_feature
 <222> (1) ... (640)
 <223> n = A,T,C or G

```
<400> 1248
aaagatataa aactatggag aaaactgcta aagggtatcc ctgaccttta tgatgatgca      60
gctatttttcg aggccaaaaa atcattttac tgggcaagaa aaacatctca ttcctttgtc      120
gtgaatatcc ttgctcaggc tctttatgaa ttattttctg ccacagatga ttccctgcat      180
caactaagaa aagcctgttt tctttatttc aaacttggtg gcgaatgtgt tgcgggtcct      240
gttgggctgc tttctgtatt gtctcctaac cctctagttt taattggaca cttctttgct      300
gttgcaatct atgccgtgta tttttgcttt aagtcagaac cttggattac aaaacctcga      360
gcccttctca gtagtggtgc tgtattgtac aaagcgtggt ctgtaaatatt tcctctaatt      420
tactcagaaa tgaagtatat ggttcattaa gcttaaaggg gaaccatttg tgaatgaata      480
tttggaaact accaagtcct aagagacttt tggaagagga tatatatagc atagtaccat      540
accacttata aagtggaaac tcttggaacca agatttggtat taatttgttt ttgaagtttt      600
tggnatataa atatgtaaat acatgcttta attgcaattt      640
```

<210> 1249
 <211> 1108
 <212> DNA
 <213> Homo sapien

<220>
 <221> misc_feature
 <222> (1) ... (1108)
 <223> n = A,T,C or G

```
<400> 1249
caaaataaat ttcaattcaa tgaaaagtaa ataacttagg gatctataaa tgacactgca      60
atgtatcttg ttccattttt aacaggaagt ctttcatgca aatgtgtgag tctcccagga      120
tgcatgaagc tccagccttt tcgtggtgac tcaatagagc aattgtacct taaaaatktg      180
caaccacctc cctgaaagtc ttctccacg ttattaagtg caatgyttat ggtaaatgta      240
gaagcatcat gatgaggacg aagagaacgc tgtcgttcag gggagtattt tactacaaaa      300
ttcagtagtg caaatccctt cgtataatag cctgcaaaga ctttcagtgt aactgggtgca      360
atgaactccc ggataaaatg aagccataca ttctccagat caacttgctt catgtggata      420
tcatcagttg ggacattttc ataaccacca gatatacggc tatcatgatg tttttcccca      480
gaccattttg cgtaatgttc catttcttct accaattcat cacaggncct tttcagaaaa      540
tatggggaac cmaaaagaca tctggacagg gctgttcaam ctatattttc agtgaaaatc      600
tttgaataat ccmcggttta tatacttttc cttccagtcc acaggatttt caaaaatctg      660
ccagaggtca ttgttataat gggaagtatt gtaattagca gtggataata gccttccaaa      720
ttcatgtcta ttagaaatgt acataaatac accctttggg gggctgagca tttggaatgt      780
ttccggagta ggggagtcct tttccctttg taaagtcatt tctctagcat ttccggcaag      840
agccatatca ggatccagtt tatcacgaac aaaatagctc ctttcattca tctctgatcg      900
gagtgtcttt cctttaatta agtacacatt agccatatat gggacattcc atactcctac      960
tctattccct tgaacaatat ccacataatc ttcagatcgt gcatagtatc catcaggact      1020
caatgtctcc cagaaattgg accacagctt tccatgacga gttacaagag gagcaatgat      1080
ctttctgttt tgttcaatca aaattttt      1108
```

<210> 1250

<211> 567
 <212> DNA
 <213> Homo sapien

<400> 1250
 ctgaatattg aactggaagc agcacatcat taggctttat gactgggtgt gtgttgtgtg 60
 tatgtaatac ataatgttta ttgtacagat gtgtgggggt tgtgttttat gatacattac 120
 agccaaatta tttgttggtt tatggacata ctgccctttc attttttttc ttttccagt 180
 tttaggtgat ctcaaattag gaaatgcatt taaccatgta aaagatgagt gctaaagtaa 240
 gctttttagg gccctttgcc aataggtagt cattcaatct ggtattgatc ttttcacaaa 300
 taacagaact gagaaacttt tatatataac tgatgatcac ataaaacaga tttgcataaa 360
 attaccatga ttgctttatg tttatatatta acttgtatatt ttgtacaaac aagatttgtg 420
 aagatatatt tgaagtttca gtgatttaac agtcctttcca acttttcatg atttttatga 480
 gcacagactt tcaagaaaat acttgaaaat aaattacatt gccttttgtc cattaatcag 540
 caaataaaac atggccttaa ctaaaaaa 567

<210> 1251
 <211> 655
 <212> DNA
 <213> Homo sapien

<220>
 <221> misc_feature
 <222> (1)...(655)
 <223> n = A,T,C or G

<400> 1251
 gaaagaaacc aattttaatgc caccaaacat aagcctgcta tacctgggaa acaaaaaatc 60
 tcacacctaa attctagcag agtaaacgat tccaactaga atgtactgta tatccatagt 120
 gcacatttat gactttgtaa tatgtaattc ataatacagg ntttaagggtgt gtggnatgga 180
 gctaggaaaa ccnaaggagn aggaaattat nnaaaagaac tgnaggtnaa gtataaagtc 240
 atatgcctga tttcctcaaa ccttttggtt ttcctcatgg cttctggcct tatattttta 300
 tcacaaacca agatctaaca gggntccttc tagaggatta ttagataagt aacacttgat 360
 cattaagcac ggatcatgcc actcattcat gggtgntcta tgttccatga actctaatag 420
 cccaacttat acatggcact ccaaggggat gcttcagcca gaaagtaaag ggctgaaaaa 480
 gtagaacaat acaaaagccc tcgtgtgggg ggaactgnng gctcactctt acttggcctt 540
 cattcnaaac aggttgggnc tttcntgcga ngatctctca gggnggtaaa aactttntgg 600
 ntttcaacan aanaggtttg gntgaatgat tactcggcng acacctaagg gatcc 655

<210> 1252
 <211> 672
 <212> DNA
 <213> Homo sapien

<220>
 <221> misc_feature
 <222> (1)...(672)
 <223> n = A,T,C or G

<400> 1252
 aaantgcaaa aaccagaag accaataatt ctgaaacttg gcatgagtgt gccagtcag 60
 cagcttgcaa agagaggatg tgtcagttac tacaattgct gtactccttt agctgagtcc 120
 ttcaactttc tccttcttgc cagtaaatac tacgttgtaa ttcatatgac tgagatctta 180
 gtatcacagg attttttagct cccatgcctc cttcaaaatt gtttacatgg atttgtttct 240

```

attctctgtgta ggccatatctc caaacacatt cacttctaaa tccaacacaa gtgaaggacc      300
agccaggatg aaacacttca gcaatcattt tgttaaaaat aacatcctgg tcatcaagct      360
aagcataagc acctcttgta taacaattca tcttaaaagc ttaaagtaca ataataaaaa      420
taactgcttg aaaactggaa atgaaataca acagaaaaaac tgaagcatta gtaatttttg      480
caagtaacc aggtacagta catttgattt catagagggt gttttctgat gtttaaggag      540
agggtagaag gggtaggaaa acttggcaag gaagatggaa acagcacaac cagttatttt      600
gcttttaata aagtaaattg aatgacagga gtagggaggt gacaaacaca tcnatatata      660
tttttcttat gg                                     672

```

<210> 1253

<211> 644

<212> DNA

<213> Homo sapien

<220>

<221> misc_feature

<222> (1)...(644)

<223> n = A,T,C or G

```

<400> 1253
ccaaatattt gttagaaact tctggtaact tagatggctt ggaatacaag ttacatgatt      60
ttggctacag aggagtctct tcccaagaga ctgctggcat aggagcatct gctcacttgg      120
ttaacttcaa aggaacagat acagtagcag gacttgctct aattaaaaaa tattatggaa      180
cgaaagatcc tggtccaggc tattctgttc cagcagcaga acacagtacc ataacagctt      240
gggggaaaga ccatgaaaaa gatgcttttg aacatattgt aacacagttt tcatcagtgc      300
ctgtatctgt ggtcagcgat agctatgaca ttataatgc gtgtgagaaa tatggggtga      360
agatctaaga catttaatag tatcgagaag tacacagaca ccactaataa tcagacctga      420
ttctggaaac cctcttgaca ctgtgttaaa ggttttggag attttaggta agaagtttcc      480
tgttactgag aactcaaagg gttacaagtt gctgccacc ttatcttaga gttattcaag      540
gggatggagt agatattaat accttcaaaa gagattgnag anggcattgaa acaaaaaaatg      600
yggactattg aaaatattgc ctctggtctg gcggagggtt gctc                                     644

```

<210> 1254

<211> 438

<212> DNA

<213> Homo sapien

```

<400> 1254
aaagggcatt tgaggggagg attattgcta tgaatgaaaa aaatatttta gcttagacta      60
agctacctgc cttcaaaata gtttagggac caccaccata ttttattttg tttttatttt      120
tgaacatttt tctaattgatt tggagagaaa actattttaca aaaattccac atatcagtga      180
tacaattttt tgctgtcacc aattttttat aatagcagag tggcctgttc taagaaggcc      240
atatttttta agttatcttt cagggttaaca tggaaatact ataaagttgg atgtcaaact      300
ttaatatggt ttcatgtttc tctaattttt tggaaatttt gtagacttta cacctggaaa      360
aaaagatttg taaaatcacc ggaacaattg tgtgctttat tttataggta gtgggttatta      420
gtattacatc cccatttt                                     438

```

<210> 1255

<211> 519

<212> DNA

<213> Homo sapien

```

<400> 1255
caagcacagg ggagtttata gttctgatgt ctttgacatt ttccctggaa cataccaaac      60

```

cctagaaatg	tttccaagaa	cacctggaat	ttggttactc	cactgccatg	tgaccgacca	120
cattcatgct	ggaatggaaa	ccacttacac	cgttctacaa	aatgaagcat	cttctgagac	180
tcacaggaga	atatggaatg	tgatctaccc	aatcacagtc	agtgtgatta	ttttattcca	240
aatatctacc	aaggaatgac	caggagaata	agatcctccg	atgttcgcaa	tggtgtggtg	300
tcaggaggct	gcctcttaga	caatctccag	atgtactgtg	atgtgagttt	gaaaaagagt	360
tcctgaagta	ccacatctgg	gagacatgcc	actagctgag	cttcccaaaa	gtctaccaag	420
agctgaggaa	ttgtatcttc	atccttagca	caaagcacct	taaaaacagt	aaaaggagcc	480
tctatattcc	agataaatat	agcactgata	aagcgacag			519

<210> 1256

<211> 178

<212> DNA

<213> Homo sapien

<400> 1256

ccatgcagga	gttcatgatc	ctcccagtcg	gtgcagcaaa	cttcagggaa	gccatgcgca	60
ttggagcaga	ggtttaccac	aacctgaaga	atgtcatcaa	ggagaaatat	gggaaagatg	120
ccaccaatgt	gggggatgaa	ggcgggtttg	ctcccaacat	cctggagaat	aaagaagg	178

<210> 1257

<211> 255

<212> DNA

<213> Homo sapien

<400> 1257

gggtccactt	gctgccccat	cattgtatca	ccttccttca	atcttttgge	tgccactctc	60
atgtagggat	ccacggtgag	gaacaaagct	tcaagcagga	cctctccatt	ttttaagggg	120
gggagctcag	atgtcttcaa	ctcaaagtca	ctattagtag	gatagccaac	aaagtgcctc	180
ttcagggtcc	atgtcttagt	acgaaccatc	ctgaagctca	ggagcccgaa	ggttccactg	240
cctggggaag	gcggc					255

<210> 1258

<211> 630

<212> DNA

<213> Homo sapien

<400> 1258

aaaactaaaa	gcatcactgc	tgaactccag	ctcagtcttc	ccattttata	atgaggactc	60
tgaagtttat	agaggtcaag	gacttgacca	aagctttaga	tatgtagtgt	ctgtgccctt	120
ttcctctaag	tttctcctag	agaatgtggg	ggctcaggaa	cagagaaaat	aaggtgcaaa	180
aagtagaaat	gggtggtggt	tctcaaagtg	tggtccatct	gcatcctagt	gactgggggtg	240
cttgttaaaa	tgcagattgc	tgggccttat	cccaatctga	ccaaatcatc	tcaggatcta	300
ccttttgaac	aaacttgcc	aggtcaaatt	cactcttggtg	gaagtttaag	tacttcagaa	360
acaagacagc	cacagaaggt	gcacctgcta	atttggtggc	ttccagtgcc	tcattctgtaa	420
cttctggtga	aatcctgaga	tgtcttactt	tacattgttt	acatcccata	acattccaac	480
atttagaaat	tactcagagc	ttatttttct	tacttgttta	gcactaaatg	aaaatagctc	540
cctgaagtta	aggagtttat	atacagtaat	tcattgcaagt	gtgtaaatta	aacagatgac	600
tttccccctt	aatatcta	gcacagcaag				630

<210> 1259

<211> 159

<212> DNA

<213> Homo sapien

<400> 1259
 aaaattttaca gataaaggca gttcaatact gccactgaga agtacatctc ttaacatata 60
 caacttttcag gccacagttt tgaaggctcg aagtattaag ttggtttgat gaattagtcg 120
 gttggcactt acgaacacat ttattgcctt gccatcttt 159

<210> 1260
 <211> 115
 <212> DNA
 <213> Homo sapien

<400> 1260
 aaaaatacta taattttcaaa acttccaaat ttcaacagat gccagtgttc tctccttttt 60
 tcatatggga aaattttcttt caaaattatt tgacgcttgg acaaaaattc cacag 115

<210> 1261
 <211> 280
 <212> DNA
 <213> Homo sapien

<400> 1261
 aaaatattgt ttatctttat ttattttgtg gtaatatagt aagttttttt agaagacaat 60
 tttcataact tgataaatta tagttttgtt tgtagaaaaa gttgctctta aaagatgtaa 120
 atagatgaca aacgatgtaa ataattttgt aagaggcctc aaaatgttta tacgtggaaa 180
 cacacctaca tgaaaagcag aaatcggttg ctgttttget tctttttccc tcttattttt 240
 gtattgtggt catttcctat gcaaataatg gagcaaacag 280

<210> 1262
 <211> 144
 <212> DNA
 <213> Homo sapien

<400> 1262
 aaattatttg atgagttcca cttgtatcat ggctaccgc aggagaagag gagtttgtaa 60
 actgggccta tgtagtagcc tcattttacca tcgwtgtgat tactgaccac atatgcttgt 120
 cactgggaaa gaagcctgtt tcag 144

<210> 1263
 <211> 487
 <212> DNA
 <213> Homo sapien

<400> 1263
 aaacatcttg ataatttggt gttgagagct gttcattcta aaatgtaatg aaattcagtc 60
 tagttctgct gataaagatc atcagttttg aaaggttact gattttcctc ttccctctta 120
 gttttttacc caatatatgg agaagagtaa tggccaatct taacattttg ttttaattgt 180
 ttaataaagc tgctgggcag tgggtgcagca ttcctaccta gtgtcataaa agcaaaaatac 240
 ttacatagct ttcttaaaat ataggaatga cattacattt ttaggagaaa gtaagttgct 300
 ttgcaccgcc tacttaattc ttttccatat attgtgatac aaacttttga atatggaatc 360
 ttactatttg aatagaaatg tgtatgtata atatacatc atacataagc atatatgtgt 420
 gtgtgtgtgt gtatatatat atatatgcat gctgtgaaac ttgactacac aacataaatc 480
 acttttt 487

<210> 1264
 <211> 250

<212> DNA

<213> Homo sapien

<400> 1264

ctgcttcaac	agagtggcag	caaccaagct	ggagtccaag	ccccctgata	aaaggcagcc	60
aatccttctg	tctgtcatca	aacgttttctt	tacagcatta	ttaaaaagga	tcctgaggtt	120
gttcttcaca	gtttctatct	caaaacctgg	aaagagtttc	tccacattgt	catagagggc	180
gtgcaggggt	tcaccccgac	agtgatgata	tttaaccatt	tccacggatg	caactttgcc	240
atttggtttt						250

<210> 1265

<211> 394

<212> DNA

<213> Homo sapien

<400> 1265

aaatatttgt	tccaaccttt	ttcgttgggtg	gcatttatgg	ctttggagca	ctgtcaggcc	60
catgttcatt	accgtgagct	cctgtgcac	tcctaatttc	caaactagcc	tggaacgc	120
ctccattgac	catgattggt	tcattggtcct	gtgcatggaa	catcatatgt	tcaggagat	180
aaagaactct	gatagtggca	cctgggtaaa	aagtacaatc	cattatatct	ggatatcaag	240
atcttttgca	gttgaagaga	ggtattgcca	cagagaaaat	tataggagca	gaagaaagtc	300
aatgaaagtc	aatgatgaca	ctccattagg	aaccagaaag	atggtattta	tttatacata	360
taataggtgt	aagagattag	aggaagcctg	tcac			394

<210> 1266

<211> 229

<212> DNA

<213> Homo sapien

<400> 1266

ccacagttgt	atcatatagc	atctctaaca	tttcatctag	gattatctag	tatagatctt	60
actatatttg	gggctatgtt	gtatacaatg	ttaacaagaa	catatcttct	ctgcatatat	120
gtgtgaatta	taaagaaaag	catgagaatg	actctaagtt	caacaaacat	gggtgaatct	180
ctatgtgtct	ccagtgtcct	ggatgggtct	cccagcaagc	cattcctcc		229

<210> 1267

<211> 722

<212> DNA

<213> Homo sapien

<220>

<221> misc_feature

<222> (1)...(722)

<223> n = A,T,C or G

<400> 1267

aaatcttctc	aactttccaa	attttcatac	taaaatatat	tattgtatta	atacaaaacta	60
cagtattata	cactacactg	tgtaataaat	aaagaaatat	aaaaataaga	cacataaata	120
taaaagtttt	ctaaaactaa	aagtacatat	gtcagtaaga	agggatttaa	tactgccagg	180
tttgaagaca	tacagtacaa	aaatgttgca	cagatctata	aactaaaaga	aataaaaataa	240
tactgatagg	taaaaatcag	ctaattgtgt	taataaattg	gtcccataat	aactaacatt	300
tggaacagct	tatgagccaa	ataacaatag	catgtccatg	tctgaaatgc	aagtacatgg	360
ataaagcaga	ttagaaaatt	tccctttcgt	ttctgtagag	aaattctgaa	aatcaatcaa	420
cataaaatca	ataccgagga	attgaaggat	gaaatgtccc	agtgtttcag	tttctctgac	480

agagtcagtg gttttaagtt ttatttgga attttgatac aagagacaaa tcaacaaatg 540
 ctagttattg taggccacac attggatgaa ggcgggtag agccttgaaa atactgagaa 600
 atggcactta cagcacacag gtcttgctta agggcaaagg agatacaaag cttcatgnca 660
 tatccttcat atggtaccac atattcaaac accatcccaa cactgatctg atgattttgc 720
 tg 722

<210> 1268

<211> 407

<212> DNA

<213> Homo sapien

<400> 1268

gatgacacaa gcagctaata accattttctg gggtttctgcc taaccccccta attgtctgtt 60
 aaagccaatt ctctgggtgt cccagttagt ggtggctttt tttctttcca cattggcaca 120
 ttcacttctc cactcttgg catgtaagaa ataagcattt acataattgg aaaaatctgg 180
 atttctgatg ccaaagggtt aaagcttctt ggatttcatt tcattgatat acagccacta 240
 ttttattttt gatcagtggc ctttgggcca ctgttcaggg tactgaccat cagtgtcagc 300
 attaggggtt tggtttttgt ttcttttggg tattttcttt ttggcacatg tgaatcttgt 360
 tttgtgtaaa atgaaattac tttctcttgt tctctgatga tgggttt 407

<210> 1269

<211> 675

<212> DNA

<213> Homo sapien

<220>

<221> misc_feature

<222> (1)...(675)

<223> n = A,T,C or G

<400> 1269

ctgaaaaaga gtgatectca atatcctaac taactgggcc tcaactcaag cagagtttct 60
 tcaactctggc actgtgatca tgaaacttag tagaggggat tgtgtgtatt ttatacaaat 120
 ttaataacaat gtcttacatt gataaaattc ttaaagagca aaactgcatt ttatttctgc 180
 atccacattc caatcatatt agaactaaga tatttatcta tgaagatata aatgggtgcag 240
 agagactttc atctgtggat tgcgttgttt cttaggggtc ctgactga tgccctgcaca 300
 agcatgtgat atgtgaaata aaatggattc ttctatagct aaatgagttc cctctgggga 360
 gaggttctggt actgcaatca caatgccaga tgggtgttat gggctatttg tgtaagtaag 420
 tggtaagatg ctatgaagta agtgtgtttg tttcatctt atggaaaactc ttgatgcatg 480
 tgcttttcta tgggaataaat tttggtgcaa tatgatgtca ttcaactttg cattgaattg 540
 aaattttggg tggatttata tgtattatac cctgtcacgc ttctagttgc ttcaaccatt 600
 tataccattt tgnacatatt tttacttgna aatatttacc tgncccggcc ggccgtcgaa 660
 agggcgaaat tcaac 675

<210> 1270

<211> 268

<212> DNA

<213> Homo sapien

<400> 1270

ccatcctggg cggagctaaa gttgcagaca agatccagct catcaataat atgctggaca 60
 aagtcaatga gatgattatt ggtgggtggaa tggtttttac cttccttaag gtgctcaaca 120
 acatggagat tggcacttct ctgtttgatg aagagggagc caagattgtc aaagacctaa 180
 tgtccaaagc tgagaagaat ggtgtgaaga ttaccttgcc tgttgacttt gtcactgctg 240

acaagtttga tgagaatgcc aagactgg

268

<210> 1271

<211> 307

<212> DNA

<213> Homo sapien

<400> 1271

cctactcttc	tccgtccatt	gtactatctg	cccgtggtgg	ggatggcagt	aggatcatat	60
ttgatgactt	ccgagaagca	tattattggc	ttcgtcataa	tactccagag	gatgcgaagg	120
tcatgtcctg	gtgggattat	ggctatcaga	ttacagctat	ggcaaaccga	acaatttttag	180
tggacaataa	cacatggaat	aatacccata	tttctcgagt	agggcaggca	atggcgtcca	240
cagaggaaaa	agcctatgag	atcatgaggg	agctcgatgt	cagctatgtg	ctggtcattt	300
ttggagg						307

<210> 1272

<211> 798

<212> DNA

<213> Homo sapien

<220>

<221> misc_feature

<222> (1)...(798)

<223> n = A,T,C or G

<400> 1272

ccattgctag	aaattgaatc	acaaataata	gctaataatt	tttcattttt	caaaaaagat	60
catttgata	gcagctatgt	ataaaatgga	aaataaaaaa	ttattctatt	ttgcatgaat	120
agttcagact	ttcccatacc	acagccaagc	agtaactaaa	attaggatct	taattttcaa	180
tgataaaaagg	tctaaggttc	atttaattat	gctcctttta	cactgtcttt	ctagattttt	240
caccagat	tttcaaaatt	tgggaatgta	aacaattgat	atattttattg	tatggtggct	300
agcagttcat	ccttctgcaa	aatatgcatt	cagagaaatg	tgaagcttgt	tttaatgaag	360
acttaaacca	tttgtgtcat	tttgtttttc	atattcaaat	acaccaaatt	aaaattctga	420
acctatattt	ttcatcatta	acttccta	ataccagaac	atataccttt	ttcatgtaaa	480
gttggaatg	ggatattgga	gttttatttt	tgaaaaatat	gtaacatgac	tttaatat	540
ttatagtttt	cagaattaga	aacataggaa	gggaaaatgt	tttaattaga	taagtcaact	600
ttttatgggc	tnagtggnng	actataatag	caaattataa	agcattatta	aatgggtata	660
ataattttta	tattacctca	ttatgaatta	actaaaataa	agnngagtga	tattttta	720
gggtgntcat	actggagctc	ctgagatata	tgatttgcta	ttgactcact	ggntgattga	780
ataatatatt	actcgcgg					798

<210> 1273

<211> 664

<212> DNA

<213> Homo sapien

<220>

<221> misc_feature

<222> (1)...(664)

<223> n = A,T,C or G

<400> 1273

aaaatatacc	ttttcacagg	tagcaagaaa	tagtacatgt	aataagtctt	tatgactgga	60
atgatccaga	aatatcacia	agcatgagta	aacacatata	taaaagtagc	tcatcatttc	120

```
<210> 1274
<211> 153
<212> DNA
<213> Homo sapien
```

```
<210> 1275
<211> 504
<212> DNA
<213> Homo sapien
```

```
<210> 1276
<211> 533
<212> DNA
<213> Homo sapien
```

<210> 1277

<211> 78
 <212> DNA
 <213> Homo sapien

<400> 1277
 ccacaggaag ttgcaaaaat tagatggact ctgtgtagct agccactctt gagtgtcagg 60
 tctgcatatg tgagtttt 78

<210> 1278
 <211> 560
 <212> DNA
 <213> Homo sapien

<400> 1278
 aaatatctaa aacaatggcc cactgaagaa aggaacaatt aactctttaa ttaattcctt 60
 aggataagta ccagaaaatt taacagctag ggcagacttc taatacaata ccgaaagtcc 120
 ttccaaaaac caagtgggtg ccaacttatg tcccttagca ttataacatt cttgagccaa 180
 tagtgtaaaa atacgctgac aatttttatg gcaaacatta ctcaagggtat cttactttcc 240
 acttattact aaagtaatta acccctaaat agatgctcct caacagtggg actacatcct 300
 ggtaaaccta tcataagttg aaactatcaa gttgaaatgc atttagtacc cggataaacc 360
 tatcataaag ttgaaaattt gttaaattgaa ccagtgtaaa tcagaggcca tcttacttca 420
 tactcatgaa gcaactatag tgggatattt ttcaacttac gagatagcct aggcttggtg 480
 aaacactgtc ctaattttact ggctctcttg taattaagtc ataaatggtc aaacatcaaa 540
 ttctagaaaa gcatatattt 560

<210> 1279
 <211> 580
 <212> DNA
 <213> Homo sapien

<400> 1279
 aaaggagatt gtttcaaaaat atttttgcaa attgagataa ggacagaaaag attgagaaac 60
 attgtatatt ttgcaaaaac aagatgtttg tagctgtttc agagagagta cggatatattt 120
 atggtaattt tatccactag caaatcttga tttagtttga tagtggtgtg aattttattt 180
 tgaaggataa gaccatggga aaattgtggt aaagactgtt tgtacccttc atgaaataat 240
 tctgaagttg ccactcagttt tactaatctt ctgtgaaatg catagatatg cgcagtgttca 300
 actttttatt gtggtcttat aattaaatgt aaaattgaaa attcatttgc tgtttcaaag 360
 tgtgatattt ttcaaatag cttttttata gtcagtaatt cagaataatc aagttcatat 420
 ggataaatgc atttttattt cctattttct tagggagtgc tacaaatgtt tgtcacttaa 480
 atttcaagtt tctgttttaa tagttaactg actatagatt gttttctatg ccatgtatgt 540
 gccacttctg agagtagtaa atgactcttt gctacatttt 580

<210> 1280
 <211> 307
 <212> DNA
 <213> Homo sapien

<400> 1280
 aaacacatac gaagaaatca actgtgatta tgaagtggca gccagctaaa tatgtcttgt 60
 atttgtctct ttcctttttt tgccctaactc atcctttact tccattcctg cttccatggt 120
 aatgcaggct caaataaatt actaggatac aagattactt caagcctctt ttctgtggaa 180
 ctcataatat gataagcatt tgttacaaga ttgctgttag ttgtttaggg gataaattat 240
 attagggaaa gaaagtcttt ctttagtttg tttaaatttc tattataatt ggttactaaa 300
 tttattt 307

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<220>  
<221> misc_feature  
<222> (1) ... (745)
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<400>	1284									
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atttacacca	agaacttctc	aataaaaagg	aatcatgaat	gctccacaat	ttcaacatac					120
cacaagagaa	gttaattttc	taacattgtg	ttctatgatt	atttgtaaga	ccttcaccaa					180
gttctgatat	cttttaaaga	catagttaa	aattgctttt	gaaaatctgt	attcttgaaa					240
atatccttgt	tgtgtattag	gtttttaaat	accagctaaa	ggattacctc	actgagtcac					300
cagtaccctc	ctattcacgt	ccccaaagat	atgtgttttt	gcttaccccta	agagaggttt					360
tcttcttatt	tttagataat	tcaagtgcct	agataaaatta	tgttttcttt	aagtgttttat					420
ggtaaaactct	tttaaagaaa	atttaatatg	ttatagctga	atcttttttg	taactttaaa					480
tctttatcat	agactctgta	catatgttca	aattagctgc	ttgcctgatg	tgtgtatcat					540
cgggtgggatg	acagaacaaa	catattttat	atcatgaata	atgtgctttg	taaaaagatt					600
tcaagttatt	aggaagcata	ctctgttttt	taatcatgta	taattattcca	tgatactttt					660
atagaacaat	tctggcttca	ggaaagtcta	gaagcaatat	ttcttc aaat	aaaanggggt					720
taaactttaa	aaaaaaaaaa	aaaaa								745

<400>	1285								
cgacggtatc	gataagcttg	atatcgaatt	cctgcagccc	gggggatcca	ctagtтата				60
atagtaatca	attacggggt	cattagttca	tagcccatat	atggagttcc	gcgttacata				120
acttacgcta	aatggccgcc	accgcggttg	agctccagct	tttgttccct	ttagtgaggg				180
ttaattqgc									190

```
<400> 1286
ctgcatcttt ctacaattct accagcaata tatgagggtt acaattttct yccatctttg      60
tgaacgcttg ttagagtctg tctcttttct ttccattctg tggggttggt ttttactttc      120
taaatggtag aaccttcaaa gcacaaaggt ttt                                     153
```

<400> 1287						
aaaaacacaa	aacactagaa	cagttgctat	gaaattactg	ataatgatcc	ctttaataaaa	60
ctgcaattaa	ccactaatat	agaaattcaa	tttaagcaag	aagttttata	tattatactt	120
tacagaaaaa	aataattttg	aaaagaataa	gmcaaacaga	gatcaaacat	ttagggcatt	180
agttactqca	ttctcttttt	agaatataca	ttaagtaaca	ctagtaaaaat	tt	232

```
<210> 1288
<211> 90
<212> DNA
<213> Homo sapien
```

<400> 1288
 aaacttagtg actatthtagt tcaattgytc atccattttt tatttgcttt tataattgcc 60
 tccttgtttt ggtatattgt aaaataattt 90

<210> 1289
 <211> 670
 <212> DNA
 <213> Homo sapien

<400> 1289
 aaatcacaaa gtaaggcacc attggattaa acattttctcc tggctttttac taagtaaaat 60
 gcatagttaa ataaatactg aacactgagt tttaatactg taatacattt caatataaaa 120
 taagaggtga atgttaaaat actgtattac atgttgaata cttttatctg aaaatgttat 180
 aaaaaaacac acatgtaagc tctgatttca gggaagaaaa attcattttt gtaattttcc 240
 atagttaaag attttaccac agaacttatt catagtttta gatgcaatta ggttgcaaac 300
 tttcaaagaa aggggtgtagg tgtattaatg aaacagtcac ttaaacta ctttctaaaa 360
 caatctattc tggatgaatg gcaactttga gctatcacc cgtttcagat ttagaacggg 420
 acctgccaag ttcagatatg caaaggaatt gtccaattct tactaccctt tataaaattc 480
 agactcactt tctctgagtc agacttttct cgtcatatt ttctaggaag ggcaaattcc 540
 atcttttgtg aaatgggtca ttaggcttta tcatagggat gtttttctact gttgaaatca 600
 gataaaagaa tcccaaataa atgatgctgc taaattacca aactgctaga gattaaaaaa 660
 attttttttt 670

<210> 1290
 <211> 352
 <212> DNA
 <213> Homo sapien

<400> 1290
 aaacaatgct acaccattt ttggcaaagt gctgtattgt tcagtctgtg tacaaaactg 60
 accatctatg aaccaatcag tataaaaaat ttctataaaa acaaaattta gacagtggct 120
 caagaaaaca agctgccatt tatgcataga ttgatgtaca gtaacctaac caaatgtccc 180
 ttttgaattt tcaagttact gaaaaaaaaat gtgtcgagaa acacattaag aaggcacatg 240
 tacagtctac aatactcttc agtctcccta actcatgccc tgcctctata aaggaaatat 300
 gttcacaaat ttacttgaga aaaaaaaaca aagccactta aaaaaaaaaa aa 352

<210> 1291
 <211> 99
 <212> DNA
 <213> Homo sapien

<400> 1291
 aaaaattatt taaggtaatg gtgttacgaa tggtttaaaa atgtctggtg acttgcttat 60
 ttttaagtga tcaccattaa gtcagaaaaa tgtattttt 99

<210> 1292
 <211> 295
 <212> DNA
 <213> Homo sapien

<400> 1292
 aaatatacct ttattttctca aactcaaagc tttatcaagt tctaacacat tttgcattga 60
 caagtgtatt tatctgcac cagtaagggt agtgaccacc acgaaagagg aatccccaga 120
 cctcctaggg actaagaaat atttcaaagg ctatgcaaat atagaacaaa aagcttttcaa 180

```
<210> 1293
<211> 256
<212> DNA
<213> Homo sapien
```

```
<210> 1294
<211> 90
<212> DNA
<213> Homo sapien
```

```
<210> 1295
<211> 519
<212> DNA
<213> Homo sapien
```

```
<210> 1296
<211> 419
<212> DNA
<213> Homo sapien
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<400>	1296						
aaagcaaac	gcagaaacca	gaagcttctg	accctctaac	atgtattact	gtccaaccca		60
ccatgagaag	tatgttcact	tggtgacaac	aaagagactc	cgtatcatat	gtatgttaat		120
gaccagattg	ttcatatggg	atTTTTctta	acagattatc	aggttgagaa	tgattctttt		180
tctccaaggg	caagaaaaag	ctggctaaat	gctagttaat	taaatccatt	ctcaattttg		240
aactgtagag	agaacctga	cttgaatgag	atTTTTctaaa	ggaagacatt	tcttgctcaa		300
cctcaggtat	aattagatta	taaggaatct	cacgtccaga	attttatctg	ctgattgtta		360
gtatggtagg	taattggcct	taggacacta	tttctactag	aaccttttac	attattttt		419

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<220>
<221> misc_feature
<222> (1)...(182)
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<210> 1305

```
<210> 1308
<211> 304
<212> DNA
<213> Homo sapien
```

<400> 1308
 ctgtcttttg gaggacgtac gtaataaggt tttaatttag taaaccaatc ctatgcatag 60
 tttcagcaact agccaaacct caccaactcc tagttctaga aaaacaggca cttggcagcc 120
 ttgtgatgtc atacagagaa gtcacaggca gtacctgagg gtctgtaggt tgcacacttt 180
 ggtaccagat aacttttttt ttctttataa gaaagcctga gtactccaca ctgcacaata 240
 actcctccca gggttttaac tttgttttat tttcaaaacc aggtccaatg agcttttctga 300
 gcag 304

<210> 1309
 <211> 289
 <212> DNA
 <213> Homo sapien

<400> 1309
 gggatttcca attaacagta ttaccagata aatattcttg gtccaagcag aaaatatcaa 60
 caaaaagagc cttcttctcc tgtaaatctt aaatgcctac atcactcttt atgatacatg 120
 gatcatctta tgtggatact taaatttttc atgtctgctt cttttgcctc tcccaactat 180
 actatgagga aattcggaac aaagacattt ttgtaatat tcttatctcc ttcacaccta 240
 gtatagagct gattttacaa aggcatttaa gagatatttg aattgattt 289

<210> 1310
 <211> 534
 <212> DNA
 <213> Homo sapien

<220>
 <221> misc_feature
 <222> (1)...(534)
 <223> n = A,T,C or G

<400> 1310
 tgctttgcat tttctgatgt attacatgac tgtttctttt gttaaagagaa tcaactaggt 60
 atttaagact gataatttta caattttatat gcttcacata gcatgtcaac ttttgactaa 120
 gaattttggt ttactttttt aacatgtggt aaacagagaa aggggccatg aaggaaagtg 180
 tatgagttgc atttgtaaaa atgagacttt ttcagtggaa ctctaaacct tgtgatgact 240
 actaacaat gtaaaattat gagtgattaa gaaaacattg ctttgtggtt atcactttaa 300
 gytttgacac ctagattata gtcttagtaa tagcatccac tggaaaagggt gaaaatgttt 360
 tattcagcat ttaacttaca tttgtacttt agagtatttt tgtataaaaat ccatagattt 420
 attttacatt tagagtattt acactattga taaagtttgt aaataatttt ctaagacagn 480
 ttttatatan gctacagggt gccctgattt tcttattgaa tttgggttaga ctag 534

<210> 1311
 <211> 114
 <212> DNA
 <213> Homo sapien

<400> 1311
 aaaatttgta ggagttgtag actacctaaa tttttaagtt atggyatttg gtcataaggtt 60
 gactgggtag gtaaagaagg aaacagacaa gaaaatggct tcttgagggtg gcag 114

<210> 1312
 <211> 95
 <212> DNA

<213> Homo sapien

<400> 1312

gggcgggtaa	aggtaggccg	cgagagcgag	gttaggagag	gataggaggc	cgcagtactg	60
ctcacacgct	ccgctcttct	cccactctcg	actct			95

<210> 1313

<211> 519

<212> DNA

<213> Homo sapien

<400> 1313

aaatgataca	gtatttttagg	tatgatttaa	gactatgatt	tacctataca	ttatatatat	60
tttataaaga	tactaaacca	gcataccctt	actctgccag	agtagtgaag	ctaattaaac	120
acgttttggt	tctgaataaa	ttgaactaaa	tccaaactat	ttcctaaaat	cacaggacat	180
taaggaccaa	tagcatctgt	gccagagatg	tactgttatt	agctgggaag	accaattcta	240
acagcaaata	acagtctgag	actcctcata	cctcagtggt	tagaagcatg	tctctcttga	300
gctacagtag	aggggaagg	attgttgtgt	agtcaagtca	ccatgctgaa	tgtacactga	360
ttcctttatg	atgactgctt	aactccccac	tgctgtccc	agagaggctt	tccaatgtag	420
ctcagtaatt	cctgttactt	tacagacagg	aaagttccag	aaactttaag	aacaaactct	480
gaaagaccta	tgagcaaata	ggctgaatac	ttttttttt			519

<210> 1314

<211> 518

<212> DNA

<213> Homo sapien

<220>

<221> misc_feature

<222> (1)...(518)

<223> n = A,T,C or G

<400> 1314

ccatgggtggg	tgaagacgct	gatctgccct	gtcacctggg	gttttttatg	agtgcagaga	60
ccaggagct	gaggaaaccc	gagytccagc	ctaaggcagg	tggngaactg	gtatgcagat	120
ggaaaggaag	tggaagacag	gcagagtgcg	ccgtatcgag	ggagaacttc	gattctgcgg	180
gatggcatca	ctgcaggga	ggctgctctc	cgaatacaca	acgtcacagc	ctctgacagt	240
ggaaagnact	tgtgttattt	ccaagatggn	gacttctacg	aaaaagccct	ggtggagctg	300
aaggttgcag	gtgagcctcc	aggttttgnt	ctgagaacac	ttctctgtag	gatctanagc	360
agatgcagag	tccctcttcc	aaaagtactg	cagacactcc	tggctgctca	ctagcaatng	420
tctgcactgc	ctcccaactn	agcttctctg	caacccttaa	gaaagacaca	ttctttcttt	480
agaaagaatt	cctgctgnac	cttacatgcc	gaagtaaa			518

<210> 1315

<211> 360

<212> DNA

<213> Homo sapien

<400> 1315

tctgtgcac	caatttatta	tagwtttgta	agtaacaata	tgtaatcaaa	cttctaggtg	60
acttgagagt	ggaacctcct	atatcattat	ttagcaccgt	ttgtgacagt	aaccatttca	120
gtgtattgtt	tattatacca	cttatatcaa	cttatttttc	accagkataa	watcttratt	180
tytacgacct	atcattctga	atcaagmaca	ctgtatgttc	agtaggttga	actatgaaca	240
ctgtcatcaa	tgttcagttc	aaaagcctga	aagtttagat	ctagaagctg	gtaaaaatga	300

caatatcaat cacattaggg gaaccattgt tgtcttcact taatccattt agcactattt 360

<210> 1316

<211> 277

<212> DNA

<213> Homo sapien

<400> 1316

aaaaaacacg	tttgttatta	ccaaawagag	acggcttttag	gtaaaaataa	taaaaaccct	60
ttgcttgyat	tacytatgca	ratagttsta	tttatctggw	cwacgggyta	aaggyacagy	120
actataggwc	tctggcttga	gtmtttacgt	tcattttctta	ttgctggaat	ktcatatttc	180
ttcttgttgg	atgactaaac	cggatgatgg	tagagatggg	aagccggcat	ttactcagcc	240
ccgccctgct	cagcctcggg	agcggacgaa	ttctcag			277

<210> 1317

<211> 716

<212> DNA

<213> Homo sapien

<400> 1317

aaaatgttct	cttgagacta	gtaggcatag	aagaaagcag	aaggaaaata	aatagaaaga	60
aggtcttcta	ccttcatggc	tattcaggct	caggagggtg	gagagaaaaa	gaaggaggac	120
aaatgaacaa	gacagatgag	ggagacatcc	tctctgatat	aagatacagt	cctctctggt	180
ggatggagtc	caatttgtgt	aacttcctat	gtattttcct	agataggacc	accactattt	240
gagaaaatat	ctcactggta	acctaaagcc	aaggataata	aaccttgata	tacttaacat	300
tcaatttctt	tccagcaatg	tgataaataa	atctatcttg	tgtttctctt	gcagattgta	360
aaagcattag	aacatttaca	tagtaagctg	tctgtcattc	acagaggtaa	gcattccatga	420
gctgccttgg	ctgttccttt	gataaagttc	atctctttca	cctggagtcc	gtctctaccc	480
ccagtccccc	atgggtggaa	gtagaattga	ctcaggcaag	agaactaagg	ggcttttcctt	540
tgagattgga	tagcaaacca	tataagtagt	attccttatc	atggctgagg	acataagaag	600
aagacgtgat	ctttgtctta	catccaaatt	gaatataaac	acttggtagc	aagcagagct	660
atgagatcat	atcattgaga	attttagaga	atatgataaa	aattgatctt	gtctgg	716

<210> 1318

<211> 515

<212> DNA

<213> Homo sapien

<400> 1318

aaagctgtat	catgttgagt	aaacctgacc	tgagccagcg	gtttaaggcg	atthttgctcg	60
atgaaggcca	agacgtgaac	ccggctcattg	ccgacttggt	aaggatacag	cgcactctgca	120
aagtaaccgt	cggcgaccct	caccagcaga	tttaccgttt	ccgtggtgcc	gaagacgctc	180
tcaacagcga	ttggatggcc	gatgcagagc	gtcactacct	gacccagagc	tttcgcttcg	240
gtccagcagt	cgcgcagtgtg	gctaacatca	tactttttta	caagggtgaa	actcgaaagc	300
tgcaagggtt	aggccccaac	acctcaggtta	aacgtgcgct	tcctgaagac	ctaccgcac	360
gcacatacat	ccatcgacag	gttaccggcg	tcataagagaa	cgcgcttagc	ttggtagcga	420
gcaatccaaa	gatctattgg	gtagggtggca	tcgacagtta	ttcattgcgc	gacctggaag	480
acttgatatc	gttcagccgc	aaccaaaacc	aagcc			515

<210> 1319

<211> 141

<212> DNA

<213> Homo sapien

<400> 1319
aaatttagtg tctcatttgg aaataaactc tgggcctatt agttgttgag tattttttttt 60
ttttactacc taaaaaaaga tttgttaaga gctgaattac aacttagcat tacataatat 120
aaaacactgt aatgtgtatt t 141

<210> 1320
<211> 497
<212> DNA
<213> Homo sapien

<400> 1320
aaattcagtc ctaagaaaga ggagtgcttg tcccctaagg gtgtttaatg gcaaggcagc 60
cctgtctgaa ggacacttcc tgcctaaggg agagtggtat ttgcagacta gaattctagt 120
gctgctgaag atgaatcaat gggaaatact actcctgtaa ttccctacctc cctgcaacca 180
actacaacca agctctctgc atctactccc aagtatgggg ttcaagagag taatggggtt 240
cataatttctt atcaccacag taagtctcta ctaggcaaaa tgagagggca gtgtttcctt 300
tttggctactt attactgcta agtatttccc agcacatgaa accttatttt ttcccaaagc 360
cagaaccaga tgagtaaagg agtaagaacc ttgcctgaac atccttcctt cccacccatc 420
gctgtgtgtt agttcccaac atcgaatgtg tacaacttaa gttggtcctt tacactcagg 480
ctttcactat ttccttt 497

<210> 1321
<211> 344
<212> DNA
<213> Homo sapien

<400> 1321
ctgtccaatg acaacaggac cctcactcta ctcaagtgtca caaggaatga tgtaggaccc 60
tatgagtgtg gaatccagaa cgaattaagt gttgaccaca ggcacccagt catcctgaat 120
gtcctctatg gccagacga cccaccatt tccccctcat acacctatta ccgtccaggg 180
gtgaacctca gcctctcctg ccatgcagcc tctaaccacac ctgcacagta ttcttggtctg 240
attgatggga acatccagca acacacacaa gagctcttta tctccaacat cactgagaag 300
aacagcggac tctatacctg ccaggccaat aactcagcca gtgg 344

<210> 1322
<211> 110
<212> DNA
<213> Homo sapien

<400> 1322
ccaccacata gccagccagg aatcccttga ggaacgggga ggacaacagc gagccacctt 60
ggcccactcc actgttgact tcgtcttcta cacgccgctg caggctttcc 110

<210> 1323
<211> 359
<212> DNA
<213> Homo sapien

<400> 1323
ccacgtgtgt ggcttgggct ggcgtctcct gctgtgagct ggctgaggag gacttctctg 60
cggctctccc cttagatccg cgctatcgtg aggtccacta tgcctgtctg gatccttctt 120
gcagtggctc ggggtgagatg gtgagaaggc gtggctgagg gactcagagg tccacagcag 180
cttagacctg gagtcatctg ttttggtctt agttctgaca ctttaatggg cttgggaccc 240
tgagacaaaa gttctcctct gtgaagcgag gatttcagga gcgaggattt caggactgag 300

gcagcctgtg aagctgtgta accgagacac gcttttcctt aggtatgccg agcagacag 359

<210> 1324

<211> 258

<212> DNA

<213> Homo sapien

<400> 1324

caatcacaca	accacaaaaa	agatactgtg	tgctctcact	ttccaaaatt	ctgcctggtc	60
tmctcctgag	gaaagyagtg	atatggtagc	tggtgtggat	cccctaaagg	aattataaga	120
tggartgyga	rgaacattat	cttagactat	aakactgkct	gcatrcrgat	atgktstcra	180
agattattcc	tgctgcraat	aaagakmttg	skaaagagca	rtatasagct	atcacagtct	240
attgacccam	asatgttt					258

<210> 1325

<211> 534

<212> DNA

<213> Homo sapien

<400> 1325

ctgtccaatg	gcaacaggac	cctcactcta	ttcaatgtca	caagaaatga	cacagcaagc	60
tacaaatgtg	aaaccagaa	cccagtgagt	gccaggcgca	gtgattcagt	catcctgaat	120
gtcctctatg	gcccggatgc	ccccaccatt	tcccctctaa	acacatctta	cagatcaggg	180
gaaaatctga	acctctcctg	ccacgcagcc	tctaaccac	ctgcacagta	ctcttggttt	240
gtcaatggga	ctttccagca	atccacccaa	gagctcttta	tccccaacat	caactgtgaat	300
aatagtggat	cctatacgtg	ccaagcccat	aactcagaca	ctggcctcaa	taggaccaca	360
gtcacgacga	tcacagtcta	tgacagagcca	cccaaaccct	tcacaccag	caacaactcc	420
aaccccgtag	aggatgagga	tgctgtagcc	ttaacctgtg	aacctgagat	tcagaacaca	480
acctacctgt	ggtgggtaaa	taatcagagc	ctcccgggtca	gtcccaggct	gcag	534

<210> 1326

<211> 177

<212> DNA

<213> Homo sapien

<400> 1326

ctgcattatg	tgtgtttaga	acgagaagtt	gtttgtacag	tattttttcta	ttgaccgctt	60
ccgtcttgcc	tgaaacctgg	gcattctttc	caatagacag	aaaatcagag	agtcaaactc	120
gatgogcaat	gagttgttct	gagaccagta	atccacgggtg	ctgcaatttg	ggtttttt	177

<210> 1327

<211> 266

<212> DNA

<213> Homo sapien

<400> 1327

aaacttgttt	tatctaatac	tgagcactgt	ttttttgtca	agtattttttt	taagaccaca	60
taattctttt	tgtctgctca	aggaaaggat	agataaataa	ttggcacaca	tttgtttctc	120
actgaatttt	acagtagtaa	attaatgtta	taatgtacca	catggagatg	agttggtaag	180
aatcatctta	gttccagagc	ccagggatta	taaacagtag	gtgaaataga	tttatgactt	240
acgaaatatg	ttgtgacaat	atattt				266

<210> 1328

<211> 409

<212> DNA
<213> Homo sapien

<400> 1328
ctgtccaatg gcaacaggac cctcactcta ttcaatgtca caagaaatga cgcaagagcc 60
tatgtatgtg gaatccagaa ctcaagttagt gcaaaccgca gtgaccagc caccctggat 120
gtcctctatg ggccggacac ccccatcatt tccccccag actcgtctta cttttcgga 180
gcgaacctca acctctcctg ccactcggcc tctaaccat ccccgagta ttcttggcgt 240
atcaatggga taccgcagca acacacacaa gttctcttta tcgccaaaat cagccaaat 300
aataacggga cctatgcctg ttttgtctct aacttggcta ctggccgcaa taatcccata 360
gtcaagagca tcacagtctc tgcactctgga acttctcctg gtctctcag 409

<210> 1329
<211> 136
<212> DNA
<213> Homo sapien

<400> 1329
ccattttcgc acagtccacc ataaaattga aaagattgac cagagacaga tcatggaggg 60
cttggcaatc tgtactgatg aagccatgga ccagaagaga agtgagtcaa tgaagagagt 120
ttctcttttc acatgg 136

<210> 1330
<211> 311
<212> DNA
<213> Homo sapien

<400> 1330
ctgctaacag ccctaacggt gcaacacaag tacaaaactca ggaacctctt cgactgccac 60
gcccttcacc aacagaagga agacagtggc gccaccacaa gtggcagggc acaggggctt 120
ctgtgacaac aatatgtcct tctagtatac attcattgca aaggctgccc tgaagtttcg 180
tttttgga aaactgttat catacatttt gtatgatgtt gcttgtgggc accatgaaga 240
gagcctggct gttaaaggaca gagggagcta aaccaacaat gcatggccct gcgtgcccac 300
aagagggagc c 311

<210> 1331
<211> 613
<212> DNA
<213> Homo sapien

<400> 1331
ctggggccakg agctgtgccc ggtgacctga gccttcataa gcacacacgt ccattcccta 60
ctaaggccca gacctcctgg tatctgcccc gggctccctc atcccacctc catccggagt 120
tgcccaagat gcatgtccag cataggcagg attgctcggg ggtgagaagg ttaggtccgg 180
ctcagactga ataagaagag ataaaatttg ccttaaaact tacctggcag tggctttgct 240
gcacggctctg aaaccacctg ttcccacctt cttgaccgaa atttccttgt gacacagaga 300
agggcaaagg tctgagccca gagttgacgg agggagtatt tcagggttca cttcaggggc 360
tcccaaagcg acaagatcgt tagggagaga ggcccagggg ggggactggg aatttaagga 420
gagctgggaa cggatccctt aggttcagga agcttctgtg caagctgcga ggatggcttg 480
ggccgaaggg ttgctctgcc cgccgcgcta gctgtgagct gagcaaagcc ctgggctcac 540
agcaccacca aagcctgtgg cttcagtcct gcgtctgcac cacacaatca aaaggatcgt 600
tttgttttgt ttt 613

<210> 1332

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<220>
<221> misc_feature
<222> (1)...(591)
<223> n = A,T,C or G
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<210> 1333
<211> 379
<212> DNA
<213> Homo sapien
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<210> 1334
<211> 384
<212> DNA
<213> Homo sapien
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<210> 1335
<211> 555
<212> DNA
<213> Homo sapien
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<400> 1335
aaattagttg ctataaattc atcaataactt tttttcccta ttatatatttt ggttctatta 60
ggatttactt aactgaatct tataacaatt cgagggtgaac tgtggcaatg aaaaccagaa 120
acagttaatg agatgcttca gctcacagtt tgaagtgtg agaacctaa tattttgctg 180
tacgggtactg agctgtacca aaatatgatg gtttaggttt atgtgcaaga ctttgtgttg 240
tagtctagac aaaggggtgg gcaagagaca tgcaaagctg aagccctgct tgaaaagacc 300
cttcaaggaa gtaaaatggc aggggcagag tgcagcttaa catgttgcta tccctgttgt 360
ttttgagttg gttttggaat ggattcaagt tcttacacaa tttattttga atacaagcat 420
aatctaggtg atttgagtta atgaacttct tttcatgatg tagggaaagc tgaatgtata 480
tatttctaag aagaatttgt ttagcagatt acaagttggc aaaatagact gttcacagaa 540
actaggcaaa aattt 555

<210> 1336

<211> 505

<212> DNA

<213> Homo sapien

<400> 1336
cctggaaaga agcccagcaa aaggttccag atgaagaaga aaatgaagag agtgacaacg 60
aaaaggaaac tgaaaagagt gactccgtaa cagattctgg accaaccttc aactatcttc 120
ttgatatgcc cctttggtat ttaaccaagg aaaagaaaga tgaactctgc aggctaagaa 180
atgaaaaaga acaagagctg gacacattaa aaagaaagag tccatcagat ttgtggaaag 240
aagacttggc tacattttatt gaagaattgg aggctgttga agccaaggaa aaacaagatg 300
aacaagtcgg acttccctgg aaagggggga aggcccaagg gaaaaaaca caaatggctg 360
aagttttgcc ttctccgcgt ggtcaaagag tcattccacg aataaccata gaaatgaaag 420
cagaggcaga aargaaaaat aaaaagaaaa ttaagaatga aaatactgaa ggaagccctc 480
aagaagatgg tgtggaacta gaagg 505

<210> 1337

<211> 385

<212> DNA

<213> Homo sapien

<400> 1337
ctggtgctag tcagagctaa tgacagaatt tcagttaa ataaaagaccc ccaactgagc 60
acaccatctt gaaaaaagta tacttatcaa acagctttca atcagttcaa gagagacacc 120
ttaattgggg agaggaagaa ttgcagagta gtttgaatc atgccaattc cagatcaata 180
actgcatgtc tgttcttttg tagaaatagc ttttgcttta tattaagtaa tcacatatat 240
attctctcta tttggataag gaaaccttcg ctttatttga caatgtataa tgatatactc 300
ttctaattca cctctgtgtc ttcacaataa acatgagtaa aatttagaca agtcatggta 360
aaggtcaata taattattta ttttt 385

<210> 1338

<211> 350

<212> DNA

<213> Homo sapien

<400> 1338
aaaggtgata ttacacaaaa cctcgtcttt tgttcaactt tggatccatt ggcaattcaa 60
tggcctcaat ctcccaaac tcgccaaagt actccctgat cttttcctca gtggcttcag 120
gattcagacc cccaacgaag attttcttca ccgggtcctt cttcatagcc atggcctttt 180
tagggatcaat gacacggcca tccagcctgt gtcctctctg gtctaggacc ttctccacac 240
tggctgcac tttgaacagg ataaacccaa accctcttga ccgtccagtg ttgggatcca 300
tttttattgt acagtcaacg acctctccaa atttagtaaa atagtctttt 350

<210> 1339
 <211> 443
 <212> DNA
 <213> Homo sapien

<400> 1339
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 cctgagtaat catatcagga tgcattgttaa gctgataaaa caataagatc ccaaaatgca 120
 gtagctcaaa aaaagtagaa gttaatttat ctctggggg acagctctgg ttctcaaatt 180
 ttacaggctc agaatcacct gcagggtctg tgaaagtaca gattgctgcg ctccgcccc 240
 agagtctctg atttagtagg tgtaggctg aaccaagaat ttgcctttct aacaagctcc 300
 caagtgatgc tgatgacttg taggaatgga tttacttcta ggattagact tcagctcact 360
 ctgtttgctg aactctttct aatattttct aagttggtag actcyctgct ccaggttctc 420
 aacgtgaagg aaggaacccc cag 443

<210> 1340
 <211> 273
 <212> DNA
 <213> Homo sapien

<400> 1340
 cctcaggaac aggtaggggc agcagaatag aatagcatcc atttcccaga gaaagactgc 60
 ctttacatkt cccatgcttt tagcacaaag cagcgtctgg gccactgtta ccagaggtga 120
 gtttatacat ttacaaaatg cttaaaatcc ttgggaagca agaggaagct aaacagaagg 180
 tcccatgtta actgaaggca aattcactca acctctctag taagggaccc atgggcctac 240
 agagtgttcc ctctacaatg tgcagagtgg aaa 273

<210> 1341
 <211> 561
 <212> DNA
 <213> Homo sapien

<400> 1341
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 cctactacca ggggtgttac tcccggccca ttatgaactc ctcttaagaa gacgacggct 120
 tcaggcccgg ctaactctgg caccgccgat cgaggacaag tgagagagca agtgggggtc 180
 gagactttgg ggagacggtg ttgcagagac gcaagggaga agaaatccat aacaccccca 240
 ccccaacacc gccaaagacag cagtcttcyt caccgctgc agccgttccg tcccaaacag 300
 agggccacac agatacccca cgttctatat aaggaggaaa acgggaaaga atataaagtt 360
 aaaaaaaagc ctccggtttc cactactgtg tagactcctg cttcttcaag cacctgcaga 420
 ttctgatttt ttgtgtgttg ttgttctcct ccattgctgt tgttgaggag aagtcttact 480
 taataaaaaa aaaaaatttt gtgagtgact cgggtgtaaaa ccatgtagtt ttaacagaac 540
 cagagggttg tactattgtt t 561

<210> 1342
 <211> 159
 <212> DNA
 <213> Homo sapien

<400> 1342
 aaagatggca aggcaataaa tgtgttcgta agtgccaacc gactaattca tcaaaccaac 60
 ttaatacttc agaccttcaa aactgtggcc tgaaagttgt atatgttaag agatgtactt 120
 ctcagtggca gtattgaact gcctttatct gtaaatttt 159

<210> 1343
 <211> 76
 <212> DNA
 <213> Homo sapien

<400> 1343
 aaaatgtaaa gccaatctat caccaaaaaat ggcataaatg taaacacaag ctaattttat 60
 aatccactgc tatttt 76

<210> 1344
 <211> 726
 <212> DNA
 <213> Homo sapien

<400> 1344
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 taagaaggct aacacctaaa ccacacgcag gcatcctgaa ctcagcagct ctgatccaag 120
 gtactgagtg gagacaaaagc actcggaggt ggcaagatgt tcagcaacca agtaagacac 180
 actggcaagg catcccaccc aaagggtgaga agcacaaaagc aggccttgag aaacaaacag 240
 tcatgccagg tgcagccaga catcctgcta taagccctga ccctagtacc ccgagttcat 300
 caagtgcctc ggttttgtgt ccataaagca cagagggcac tgaccacccc aaaccagaat 360
 cccaaggaat ccttatggat ggcatagggc ctcagaactg ctgcaggatc attttccttt 420
 tcaggtcgtg gctgaacttg ttcacctga agagctcact gtcataaaat gcagagaggt 480
 tgtggatgtt gatctgacga gccttatcca ccaagtcctt mtcagggacc tcaatagtgt 540
 cctgctgggc cccaaagcgg ttgcgctgat atgtcacstg ctctgccact aactgcttca 600
 gtatgaagag caacagctca ttgtgtcac gccggaatga aaggtagcgg gcaaaagtct 660
 tgcgcatgct gcgcatgacg ctgaacttct gtgtgtctat gaagstctcc akmatcayga 720
 gratgg 726

<210> 1345
 <211> 742
 <212> DNA
 <213> Homo sapien

<400> 1345
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 ttgagggcag gctgatgttt cctgaatggg cccctgggtg ttgcttgctc ctgactctcc 120
 atttcccat ctgagtggat ttggacctaa tagggcactg gagctgggtc gaatcctgac 180
 tggactactt ggcaacttta tgtctgggag caagttactt aacctcccca agcctgtgtc 240
 tgtgaaatgc gggtaaata gaatagatgt ttggcagcag ctactccttg ttgagctctc 300
 acagtgaact ctccctgcctc tgccctcctt ccccgccctc cctgggtgcct agcgtcaggt 360
 ctageccactt cctcctgggc cctctcctt tttctgtggc tggctgcctg cccgcctggc 420
 gctggacctt tcatgtaacg ggaatcagca tgtatattct ggtctggtct gtttctacac 480
 ttaattttgt ttccagtagt atttccctgt accggcagag ttcacaaaca catttgaaga 540
 ggctttttct caggattctt aaccttccaa aggaagtccc atggatgggt ttctagaagt 600
 ctataaatgc tctgaaattg tatttttctg tggaaaagca taacttttat ctgcttggtc 660
 gtgtcaaaa aaagatcatg aatggaatga attgcattga attttatgcc attgggggct 720
 taatactaaa aggatatgga ag 742

<210> 1346
 <211> 573
 <212> DNA
 <213> Homo sapien

<220>
 <221> misc_feature
 <222> (1)...(573)
 <223> n = A,T,C or G

<400> 1346
 aaatgcattk ttaacttaca gtatatttcaa cttacgatgt gtttatcasg aagtaacccc 60
 atcataagca gaggagcatc tgtattgCGT aatttgactg gcacagttaa ttaggttctg 120
 ttcagtgwtt tccgtcaaca agatgtttat tgtgtgagta aacaagttaa gccctgtgac 180
 aagctgaata agaatagtct ctccctcagca gcttatagta aacaagggtg gtaatcctta 240
 cattagtggc tagactatca aacgaaatat ataacatgta agaacactaa agacagaatt 300
 actgtggcat agagatagtt agaattgctt cagcctaaga gatgaattag gtaatgcaag 360
 gaggtgaata tggtggcctg caatatgaac aaggcagaga gctgggagag taagatgtaa 420
 gttgctaagg agggatgtgt cttgagtttg gaaaccataa agggaaatca taggtaatgc 480
 tagagtcact gatcttangg agccttgaat aacggtgatg actaaggga tctttatattt 540
 gnggggacta ttggaattaa attggccaga att 573

<210> 1347
 <211> 333
 <212> DNA
 <213> Homo sapien

<400> 1347
 cctggtttct ggtggcctct atgaatccca tgtaggggtgc agaccgtact ccatccctcc 60
 ctgtgagcac cacgtcaacg gctcccggcc cccatgcacg ggggagggag atacccccaa 120
 gtgtagcaag atctgtgagc ctggctacag cccgacctac aaacaggaca agcactacgg 180
 atacaattcc tacagcgtct ccaatagcga gaaggacatc atggccgaga tctacaaaaa 240
 cggccccgtg gagggagctt tctctgtgta ttcggacttc ctgctctaca agtcaggagt 300
 gtaccaacac gtcaccggag agatgatggg tgg 333

<210> 1348
 <211> 185
 <212> DNA
 <213> Homo sapien

<400> 1348
 aaaaaagctt gcagcaagaa aatgccagtg tgcaactggg tgactaaaga ccaaagaaaa 60
 acagttaaaa gggacagctt acttgctctc tgtctcaggt ttaacttctc acctgaaatc 120
 tctcatagcc ctaattaaac acaaacaaaa gtctcttcca tagataggct acttctcagc 180
 ttcag 185

<210> 1349
 <211> 171
 <212> DNA
 <213> Homo sapien

<400> 1349
 gcggcagcga ggggctcgga gaggtgctcg gattctcgta gctgtgccgg gacttaacca 60
 ccaccatgtc gagcaaaaga acaaagacca agaccaagaa gcgccctcag cgtgcaacat 120
 ccaatgtgtt tgctatgttt gaccagtcac agattcagga gttcaaagag g 171

<210> 1350
 <211> 400

<212> DNA

<213> Homo sapien

<400> 1350

ttgtcatatc	atatctatgt	cacctgtgta	ttctgagatt	acacacatac	ctgccaatat	60
acctgggaaa	ggttatttta	tcacagttac	acttgagttc	ttggcaggca	ggactgagga	120
agagtaattt	gaaagaagtt	ttacatccta	tttagaagaa	atcactagta	tttccttaaa	180
taacagggtta	caatagaaag	atactgcctg	gaagttatcc	tttcactttg	gttcattttt	240
agtttttctt	tatgatttac	atagctgttt	aattcatttg	cttatagtac	aatcctgcca	300
taaagtatta	aagcacaaga	tacctattat	tccttcaaca	tctgcatttt	tcaagtttta	360
tactctacat	ccacagtagc	tcagcagttc	ttgaatgttt			400

<210> 1351

<211> 309

<212> DNA

<213> Homo sapien

<400> 1351

ccaggaaagg	gcagtcctga	gggagaagac	aggattcagg	gcagtgctcc	gaagctgtgt	60
gctcacctgg	ttggctcatc	aaacctggca	accctgtggc	ctgtctgccg	gagctgactg	120
gatccactca	tcaattcttc	gtccccacta	ctaagaactg	gcatgttttg	ctgggtgtgg	180
ctctgcactt	caggaatggg	cacaacaggg	ggtagccctc	aaaagcactc	ctttttctat	240
acctcttctc	aaggccatgt	aagttgcccc	tctctacctg	gctgtggaca	aaagggtatc	300
tgctcttgg						309

<210> 1352

<211> 268

<212> DNA

<213> Homo sapien

<400> 1352

ccacttcatc	tgtgtgggaa	cgtggtcagg	ccgggtgctg	gtgtttgaca	tcccagcaaa	60
gggtcccaac	attgtactga	gcgaggagct	ggctggggcac	cagatgccaa	tcacagacat	120
tgccaccgag	cctgcccagg	gacaggattg	tgtggctgac	atgggtgacgg	cagatgactc	180
aggcttgctg	tgtgtctggc	ggtcagggcc	agaattcaca	ttattgaccc	gcattccagg	240
atttgagatt	ccgtgcccct	ctgtgcag				268

<210> 1353

<211> 620

<212> DNA

<213> Homo sapien

<220>

<221> misc_feature

<222> (1)...(620)

<223> n = A,T,C or G

<400> 1353

cctgagtaat	tattccatca	tagacaaact	tgtgaatata	gtggatgacc	ttgtggagtg	60
cgtgaaagaa	aactcatcta	aggatctaaa	aaaatcattc	aagagcccag	agcccaggct	120
ctttactcct	gaagaattct	ttagaatttt	taatagatcc	attgatgcct	tcaaggactt	180
tgtagtggca	tctgaaacta	gtgatttgtt	ggtttcttca	acattaagtc	ctgagaaaga	240
ttccagagtc	agtgtcacaa	aaccatttat	gttaccacct	gttgacagca	gctcccttag	300
gaatgacagc	agtagcagta	ataggaaggc	caaaaatctc	cctgggagact	ccagcctaca	360

ctgggcagcc atggcattgc cagcattgtt ttctcttata attggctttg cttttggagc 420
 cttatactgg aagaagagac agccaagtct tacaagggca gttgaaaata tacaaattaa 480
 tgaagaggat aatgagataa gtatgttgca agagaaagag agagagtttc aagaagtgtg 540
 attgnggctt gtatcaacac tgttactttc gtacattggc tgggaacagt catgtttgct 600
 ttcataaatg aagcagcttt 620

<210> 1354

<211> 398

<212> DNA

<213> Homo sapien

<400> 1354

aaaggattat ttttatgcaa agtattctgt ttcagcaagt gcaaatttta ttctaagttt 60
 cagagctcta tatttaattt aggtcaaagc ctttccaaaa agtaatctaa taaatccatt 120
 ctagaaaaat atatctaaag tattgcttta gaatagttgt tccactttct gctgcagtat 180
 tgctttgccca tcttctgctc tcagcaaagc tgatagtcta tgtcaattaa ataccctatg 240
 ttatgtaaag agttatttta tcctgtgggtg catgtttggg caaatatata tatagcctga 300
 taaacaactt ctattaaatc aaatatgtac cacagtgtat gtgtcttttg caagcttcca 360
 acagggatgt atcctgtatc attcattaaa catagttt 398

<210> 1355

<211> 371

<212> DNA

<213> Homo sapien

<400> 1355

ctggytcctc agtggggaact gagtcattac ctgctaaagg gtagaagagg agagagagag 60
 gccagagcct ggggatgggg cagaagggtgc agcaggaagg aaggtagag tgagaaaaat 120
 ttccaaataa ggggtgatgt gtgagtgtc agaggggtgac tgaggacatc tccagcattt 180
 ccattgagga gggaggaagg aggggccctt gggttctggg gcagatgccg gcaggggtctg 240
 gatgagatgc ccccaacctc aacctgtgtc ctctgaaaac acttcaccca gtcacactga 300
 ggagccccctc caggcccagg ggcccccca ggtaggcgta tctcagctcc tctctggaag 360
 gacccccaca g 371

<210> 1356

<211> 338

<212> DNA

<213> Homo sapien

<400> 1356

gcggcgcggg cgggcggtaaa atgtcggttc caggacctta ccaggcggcc actgggcctt 60
 cctcagcacc atccgcacct ccatacctatg aagagacagt ggctgttaac agttattacc 120
 ccacacctcc agctcccatg cctggggccaa ctacggggct tgtgacgggg cctgatggga 180
 agggcatgaa tcctccttcg tattataccc agccagcgcc catccccaat aacaatccaa 240
 ttaccgtgca gacgggtctac gtgcagcacc ccatacctt tttggaccgc cctatccaaa 300
 tgtgttgtcc ttcctgcaac aagatgatcg tgagtcag 338

<210> 1357

<211> 159

<212> DNA

<213> Homo sapien

<400> 1357

ctgggtgtgt gcctctggag tacttccccg cagctcctca ttgctcacat agtaggcaat 60

ggcgttgctc tcaaacacac agaatccatc atcacccctca aatgctggga ccttgccggc 120
aggaaatttg cggagaaatt caggggtgcg gttggtttg 159

<210> 1358
<211> 306
<212> DNA
<213> Homo sapien

<400> 1358
cctgtcagag tggcactggg agaagttcca ggaaccctga actgtaaggg ttcttcatca 60
gtgccaaacag gatgacatga aatgatgtac tcagaagtgt cctggaatgg ggcccatgag 120
atggttgtct gagagagagc ttcttgtcct gtctttttcc ttccaatcag gggctcgctc 180
ttctgattat tcttcagggc aatgacataa attgtatatt cggttcccgg ttccaggcca 240
gtaatagtag cctctgtgac accagggcgg ggccgagggg ccacttctct gggaggagac 300
ccaggc 306

<210> 1359
<211> 382
<212> DNA
<213> Homo sapien

<400> 1359
agagggagtc cagcccccaa gccttgtgag gcaactgttar gcagataggg aaaagagggg 60
tccttagatc actggttcaa ggagggatct ggtaggggca gcatttcttc tgggctggaa 120
acagaatggg gggttcaaga tggcagaacc attccattat tggagctata agcccctaga 180
attgctccat ggcctatctc ggtttccctt ggatctcatc tgctcctgaa ctgcacctgt 240
catggcaagt ccactctccg ccccatctc ccctgagcca atgtgagtc ggtgaacaaa 300
attcattggt tcccaatca tggtcgggtc aatccgtctt ctcttctctt ttcttctcca 360
ccatccagac gttcagctac ag 382

<210> 1360
<211> 365
<212> DNA
<213> Homo sapien

<400> 1360
aaaaaacctt tcaaaataaa acttagtaaa atctagaact gkttcttggc ctacttgaga 60
ggaacttcca tattttcaca gccatctccg aaagcagcag ttgctgtaaa ttaactgaga 120
cttggaatg gtgcagactg tcttggtaga gctgttctta tagcacaatt ttatctggaa 180
aataaacttg taaatgcgtg ctgtatatta atacatgtgt gccatattt atttttatta 240
tctcctgcc gtctttgctc aatgggagat gacagaccaa cttctcaacg tgatttcccc 300
atttcattga atgacattta tatgccactt atgaaaaaaaa tactgctgtg aaagaaatgt 360
acttt 365

<210> 1361
<211> 502
<212> DNA
<213> Homo sapien

<400> 1361
gaggtatgga aaaatatcaa caaggaaata ttagatttga actgctgctt cgtagcaca 60
cagcacattc tccaggatat accatatgtt aggacacaaa acgggtctca ataaattttt 120
aaaagtcaaa atcttatcaa gtatcttctc agaccacaat ggaataaaaac tggaaatcaa 180
taacaagagg aacttctgaa attgaacaga tacacggaaa tcaaactaca tgttcctgaa 240

tgaccactgt	gtctatgaag	aaattgattt	taaaaattta	aaaattcttt	gaaacaaatg	300
aaaatagaaa	cacagcatac	aaaaatgtat	agggtacaac	aaaagaagtg	ctatgaggga	360
catttatctt	aataaacacc	cacatcaata	aggtagaaag	tttttaaaca	aataacctaa	420
taaacgcac	tcaaggaact	agaaaagcaa	gaacaaatca	aacctaaaa	tagaaggaaa	480
taaatagtaa	agatcagagc	ag				502

<210> 1362

<211> 545

<212> DNA

<213> Homo sapien

<400> 1362

ctgattggat	gtctaggaat	gactgaaaga	aaccaaaca	gcctgtccac	tgctgctgtg	60
ggatggagga	ggcgtaagca	gaaacactaa	cagtatactg	acctcttagc	agaaccgctt	120
ccattctgga	gatcacggct	gctaaatcca	gcacccccac	ttcattttac	ccccagcata	180
ttgttctgta	gtcttttctt	gaaacatctt	gattgctttt	cctcggcagc	tttcaaaaaa	240
ccaaataata	atagttatcc	gtcttctact	tcattggaaga	ttgttttggt	gccctgaccc	300
tctgaagtgc	ccagttcctg	ccatctgaaa	cctcggcctg	atctgatctc	atgttggaat	360
ctgcctgtct	ttcacacagg	gctgggtctt	gtcctttaca	tgccagtttt	gcttgtgaat	420
tcttgctttt	ttcctctcat	cagccttaag	tttaggcgtt	tggtgttctc	cagtgatgta	480
gacagttccc	ttcacaagtc	acagttcttc	ccataaatga	ggcccgtga	cctctgctgg	540
acttt						545

<210> 1363

<211> 286

<212> DNA

<213> Homo sapien

<400> 1363

gggagatgca	ggatgtagac	ctcgctgagg	tgaagccttt	ggtggagaaa	ggggagacca	60
tcaccggcct	cctgcaagag	tttgatgtcc	aggagcagga	catcgagact	ttacatggct	120
ctgttcacgt	cacgctgtgt	gggactccca	agggaaaccg	gcctgtcatc	ctcacctacc	180
atgacatcgg	catgaaccac	aaaacctgct	acaacccccct	cttcaactac	gaggacatgc	240
aggagatcac	ccagcacttt	gccgtctgcc	acgtggacgc	ccctgg		286

<210> 1364

<211> 503

<212> DNA

<213> Homo sapien

<400> 1364

ccatcaggat	catgaaaaca	aactttgggt	aatgtgagca	actgcgccag	acaggacaca	60
ggttacaggg	cctgacgtca	ctaacggtaa	ctgacaatct	tggaatggac	cctactgctg	120
atgtttcaaa	aggacacaga	ggtgaactgg	tcacttctaa	ttaagaagag	ccagtggggg	180
gggggaagct	gaaaaccaa	aatccacgta	gacatacgtg	gcagtgtgaa	cgtctgtcct	240
ccccttcctt	ctcctcactt	cctctcctcc	tcctcactca	ggctggtatt	ctcctggtgt	300
gcggatgtca	gcttgccctg	cagaagggct	gccagttttt	tagatgtctt	tttgagaaac	360
gagctgccc	gatgggcact	gttcacgtgc	aggtacaggt	cctcctgggt	ggggcccgtg	420
tagccgcaat	cctcgcagac	gtagagcttg	tcctcgccgt	gcttataggc	atactgctgc	480
tgcaccccat	ggattttctt	cag				503

<210> 1365

<211> 245

<212> DNA

<400>	1365								
ctgggcggct	ccacgctcat	ccagtgggccc	taggttctga	ctgaccagcg	aacaaaaact				60
gtgacagaga	tctaggattt	cattcaggca	gtgaaacacc	tacccgggaa	acagagttgg				120
cattaggaaa	ggaaggaagg	tacatccatg	aagttaaagt	gttaggagaa	cagtcctgatt				180
aatagctgat	ctaattaata	gctgacctcc	caaatctgac	aggatagaca	ctgccacgtg				240
caagc									245

<213> Homo sapien

```
<400> 1366
aaaatcccca taaatctttt ctgtcttgag gtagttgcaa aataaatcat aacttgata      60
tcaactagag ctgaggcttt gactttttac tcattaaaac tagttgttac aggaactacc      120
tttaqatatt t                                     131
```

<213> Homo sapien

<400> 1367									
ctgtgcagtt	atatgaccat	aaaggaaatg	aaccattaaa	aatggatcta	cagccatata				60
ttctgccgtt	actcagaggc	ttaatgattt	attttcccc	tccagccctg	cctttaccag				120
gttaaattgac	agaagacctt	ctattgtacc	tattgttcaa	aaaatattac	tgttctgtgg				180
aacctgggag	agtccaattg	ataagagaaa	ctgaatcata	ctgatgaggt	gaaggatagg				240
tctgccggtg	tggggcaggg	cactctttct	cagcagccaa	gataacttat	cacacacgaa				300
gcagagagaa	tgcaccgat	gaaaatctct	ctgaactgtg	ttccttgaag	gatctcttaa				360
aaaaaaaaaa	tctgaaacat	catccattga	acaaatgaaa	ggcttatacc	tttaccatga				420
aqaaacattt									430

<213> Homo sapien

<400>		1368						
ctgggcggat	agcacogggc	atattttgga	atggatgagg	tctggcacc	tgagcagtcc		60	
agcgaggact	tggtcttagt	tgagcaattt	ggctaggagg	atagtatgca	gcacggttct		120	
gagtctgtgg	gatagcagcc	atgaagtaac	cgaaggagg	tgctggctgg	taggggttga		180	
ttacagggtt	gggaacagtc	cgtacacctg	ccattctctg	catatactgg	ttagtgaggt		240	
gagcctggcg	ctcttctttg	cgtgagcta	aagctacata	caatggcttt	gtgg		294	

<213> Homo sapien

```
<400> 1369
ctgaaggcaa tgggggactg aggaaggagg cagcagaagt aggagaggag caagaatcca      60
gaagggaaat gagaacgaca aaactgaagt gcacttcaac atcctgcagc caaaggggta      120
```

```
<210> 1370
<211> 540
<212> DNA
<213> Homo sapien
```

```
<210> 1371
<211> 142
<212> DNA
<213> Homo sapien
```

```
<210> 1372
<211> 377
<212> DNA
<213> Homo sapien
```

```
<210> 1373
<211> 504
<212> DNA
<213> Homo sapien
```

<400> 1373
ccatgctaag tttgggaacc gctggtgatg ggacatggat gcttgcaacc gaccgtgggc 60

```

ggatgtggtt gaccagatgg cagaggacga caccatccat gagggctgcc cccaggtctt 120
cgtgcagact gaccttcaat ctcatctcaa tgctctcacg aagttgttcc accagctctt 180
tctcttctct catctgctcc attttctctc ggattgtaaa ctgcgggtct atagattcca 240
aatttctctg aggtcttaga aacacagact cagaaatcaa atgaggatgt ctcagaaagg 300
agtcactttt ccagaggcag gctgcccctt aactcagccg agcagcagga accactgggg 360
ccaaagctat tttatcttcc ttaggtaaaa aaaaatcaat agaataattc ttccccgctt 420
acatgctccc accactgatg aacgcgatct tcagcaagaa gaactttgag tccctctccg 480
aagccttcag cgtggcctct gcag 504

```

<210> 1374

<211> 201

<212> DNA

<213> Homo sapien

<400> 1374

```

cctccgtaag atgcttgaca attttgactg ttttggagac aaactgtcag atgagtccat 60
cttcagtgtc tttttgtcag ttgtgggcaa gctgcgacgt ggggccaaagc ctgagggcaa 120
ggctataata gatgaatttg agcagaagct tcgggcctgt cataccagag gtttgatgg 180
aatcaaggag cttgagattg g 201

```

<210> 1375

<211> 295

<212> DNA

<213> Homo sapien

<220>

<221> misc_feature

<222> (1)...(295)

<223> n = A,T,C or G

<400> 1375

```

ctgtgaggct gnttccaagg aggaaaacaa ggaaaaaaat cgatatgtaa acatcttgcc 60
ttatgaccac tctagagtcc acctgacacc ggttgaagggt gttccagatt ctgattacat 120
caatgcttca ttcattcaacg gctaccaaga aaagaacaaa ttcattgctg cacaaggacc 180
aaaagaagaa acggtgaatg atttctggcg gatgatctgg gaacaaaaca cagccaccat 240
cgtcatgggtt accaacctga aggagagaaa ggagtgcagg tgcgcccagt actgg 295

```

<210> 1376

<211> 318

<212> DNA

<213> Homo sapien

<400> 1376

```

ccagcgctac tgtactggcc cagggcagag ttcattgtatc tcgtcttgac cagctctaca 60
ggggaggcga tgacagtggg gcagaagcct gccccaaagg cagaagtga gtaggcaagg 120
aggatcatctg tcatgaggtt ggctttcagg agggcatcct tgatgaggtc ataggtcacc 180
agctcagcac agttgacaat ggcattacga gcaacattgg gggagggtccc tttccagagg 240
ccccggaacc ctctctctcg ggcaatggtc ttgtaggcat tgacggtgct ttggtatctc 300
cgaccacctc cagcccg 318

```

<210> 1377

<211> 143

<212> DNA

<213> Homo sapien

<400> 1377
 gtggattccg ytcggggcac cgatctcgcc aagatcctga gtgacatgcg aagccaatat 60
 gaggtcatgg ccgagcagaa ccggaaggat gctgaagcct ggttcaccag ccggactgaa 120
 gaattgaacc gggaggctcgc tgg 143

<210> 1378
 <211> 98
 <212> DNA
 <213> Homo sapien

<400> 1378
 aaatattggt aataggctcg caacagcaac tatagaagta caactcaata gatggcatta 60
 aaacatattg tagtgtggat atatatTTTT tctTTTTT 98

<210> 1379
 <211> 330
 <212> DNA
 <213> Homo sapien

<400> 1379
 aaagatgttc acgttacgct ggaccaaatt aagacggctt tctccctctt gctgacgtgc 60
 cccagccgtg ataatgacca gcttggagtt tgcagttaca ttatagtctt tgccagagac 120
 aatcttttgg gttctaagga aaaggctgcc atgttggaga tccatcatct ctcccttcaa 180
 tttgtcttcg acgacatcaa caagagcaag ttcatctgcc aagtccttca ttaagatact 240
 gatggcacag gccatgccaa cagcaccaac cccaacaact gtaatcttat tctggggggg 300
 ctgttcttcc tttagaagat tataaatcag 330

<210> 1380
 <211> 269
 <212> DNA
 <213> Homo sapien

<400> 1380
 ccactcctgg aaaccactg atagatgagt ttccccatt cttctggcct ccgccacatg 60
 atcaggaagc tggacttgct cttatccaac cactcgaggt tccctttctt cctcagttcc 120
 tctaatacaa tctggatcga ctccacagga agctttcgtc gtagcttgac gttgttgaag 180
 agcgggctct cctgagcttc catcacctgc atgctggact gtttgtgcag gcggcagaag 240
 gacaggacca gcgagcacca ggcggccag 269

<210> 1381
 <211> 232
 <212> DNA
 <213> Homo sapien

<400> 1381
 aaaagagagg aaaggcagtg cagggctgga ggtcctggag ggtggcggcg ggtcgtccta 60
 actagcaggc tgaaaggctg tggaggggat gccttcactc agaggaagtt cacagccacc 120
 tgccttgga catgtacctg ttcattcttt cgtaatgtta gtattcattt tgctatcttc 180
 ctgttgccat ttccaaacag tgtcagtatg tttttgttaa atacgaacat tt 232

<210> 1382
 <211> 348
 <212> DNA

<213> Homo sapien

<400> 1382

aaacgtgcta	aagggaaagg	aatctgacat	tctgggtaaa	tcttactcaa	tctaaatcaa	60
agcttggttt	tcaggaggag	gaaggtgcga	gcgcaggcag	aggtgctgaa	tactcctctt	120
ctgattcact	tccatcatcc	tctttctctt	ggctactgcc	ctcagtgcta	agccgggtcaa	180
acccttttctg	actgtagccc	ttacggcttg	caaagaaatt	accaagggtt	aagcctccac	240
ttccctttcc	tctaaatctt	cccagtactc	ttcctgaact	cgtctcgagt	ttgtgttcag	300
aatctccaaa	ggcccttgat	tttttccacc	gaataaatat	ggcaatgg		348

<210> 1383

<211> 293

<212> DNA

<213> Homo sapien

<220>

<221> misc_feature

<222> (1)...(293)

<223> n = A,T,C or G

<400> 1383

ctgcttcaan	acctcagctt	catgggactt	gcgtctttct	tctgcagctt	ctaatttctt	60
ctgaatttcc	tccagggaaa	gatccttctt	ctttggaggg	gaaaggggga	attctggaac	120
agattctttt	gaccgagggc	tgagaatcag	ctcaaaagcc	tggcccagag	cacgcttctc	180
cagttctttc	acctggatat	cagaagaagc	catggtgaat	agaagacaag	cgacaggcag	240
tgtattctgc	acaatcaact	gggataagga	aagtcctgct	cagtccgagc	cgc	293

<210> 1384

<211> 573

<212> DNA

<213> Homo sapien

<400> 1384

ctgaagcaac	ttgggattaa	ttgcttgatt	agcttcacga	agcacagaga	taaggctcgt	60
cacttgcttt	atgttattag	gtgtaaagaa	agtgtatgct	gtgcctgttt	tggtagtcgc	120
agcagttctt	ccaattcgat	gaatataatc	ctctgaggag	ttagggtagt	cataattgat	180
gacaaatttc	acatcttcca	catctagccc	tctggaggcc	acatctgtag	caatcagaat	240
aggagctttt	ccatgtttga	attcatctag	aaccacgtca	cgtctctgtt	gactcttgct	300
accatggata	cccattggcag	gccacccatc	tctcctcatt	ttcttggtta	gctcatcaca	360
tcttcttttg	gtttccacaa	aaacaatggt	tttattctcc	ttctcactca	tgatctcttc	420
cattagacga	ataagttttt	catccttttc	tacgtcatga	cacacatcca	caatctgaag	480
aatgtttgtg	tttgactca	gttcaagtgc	accaatgttt	atatgaatat	agtctttcag	540
gaaatcttca	gcaagctgtc	ttacttcttt	tgg			573

<210> 1385

<211> 150

<212> DNA

<213> Homo sapien

<400> 1385

ccaaggccgc	tagggctcctt	acccctcagg	atcactcccc	agccctttcc	tcaggaggta	60
ccgtctctca	aggtgtgcta	gcagtgggcc	ctgcccaact	tcaggcagaa	cagggaggcc	120
cagagattac	agatcccttc	ctgtaagtgg				150

<210> 1386
 <211> 159
 <212> DNA
 <213> Homo sapien

<220>
 <221> misc_feature
 <222> (1)...(159)
 <223> n = A,T,C or G

<400> 1386	
aaatgatggt ttgggtaaga gtggaccatg agaattagct gacagcatcc cctttctctc	60
tccctgcctt ggtgggaccc tccctgtgtg accttggtca agtcctcgaa cttttgtccc	120
gtatttaaga tggagctgnt ttacctactt cataagaca	159

<210> 1387
 <211> 735
 <212> DNA
 <213> Homo sapien

<220>
 <221> misc_feature
 <222> (1)...(735)
 <223> n = A,T,C or G

<400> 1387	
ggtgnaattc gcctttgaan ggccgccggg caggtccttt ntgtstgctg aaggcagatc	60
gcttggtcca caccagctac cactcccagg cagtgcatac ccgccctggt tgcagaaatg	120
cacgtgttac tagcatctcc tgggagctga ggcagaccct gtcagttgta tttgatgcct	180
tcacacggg gcagggaaag aaagactggt cctcttccg gatgttctcc cgaaccctca	240
cggagccctg cccctggct tcagagagcc gagtctatgt ggacatcacc acctacaacc	300
aggacaacga gacattagag gtgcaccac ccccgaccac tacatatcag gacgtcatcc	360
taggcactcg gaagacctat gccatctatg acttgcttga caccgccatg atcaacaact	420
ctcgaaacct caacatccag ctcaagtgga agagaccccc agagaatgag gccccccag	480
tgcctttct gcatgccag cggtagctga gtggctatgg gctgcagaag ggggagctga	540
gcacactgct gtacaacacc caccataacc gggccttccc ggtgctgctg ctggacaccg	600
tacctggta tctgcggctg tatgtgcaca cctcaccat cacctccaag ggcaaggaga	660
acaaaccaag ttacatccac taccagcctg ccagggaccg gctgcaaccc cacctcctgg	720
agatgctgat tcaga	735

<210> 1388
 <211> 369
 <212> DNA
 <213> Homo sapien

<400> 1388	
ctggggacag cctacagggg cctccagcct gtgccagacg aggaggtgat tgaactgtat	60
gggggtaccc agcacatccc actataccag atgagtggct tctatggcaa gggtcctcc	120
attaagcagt tcatggacat cttctcgcta ccggagatgg ctctgctgtc ctgtgtggtg	180
gactactttc tgggccacag cctggagttt gaccaagcac atctctacaa ggacgtgacg	240
gacgccatcc gagacgtgca tgtgaagggc ctcatgtacc agtggatcga gcaggacatg	300
gagaagtaca tcctgagagg ggatgagacg tttgctgtcc tgagccgcct ggtggcccat	360
gggaaacag	369

<213> Homo sapien

aagatggttt	ctggcatttt	ctttttattt	gtaagggtggt	ggtaactatg	gttattggct	60
agaaatcctg	agttttcaac	tgtatatatc	tatagtttgt	aaaaagaaca	aaacaaccga	120
gacaaacct	tgatgctcct	tgctcggcgt	tgaggctgtg	gggaagatgc	cttttggggag	180
aggctgtagc	tcagggcgtg	cactgtgagg	ctggacctgt	tgactctgca	gggggcatcc	240
atttagcttc	aggttgtctt	gtttctgtat	atagtgcacat	agcattctgc	cgccatctta	300
qctgtggaca	aaaggggggtc	ag				322

<213> Homo sapien

aaatattagw	tgagacttta	caggcacata	actgttcaga	tagaaacaaa	cataacagac	60
taaaatactt	tcaaaattaa	agccatctag	aaaatggaag	taactgaaac	tgtagccatt	120
acaattcttt	ttctggtttt	gagcaaaaat	tttatctctc	tggcaaaaaca	cctttgtctg	180
atcatttgag	agacagggtt	cttgatact	gtttcttcaa	cgtaaacctc	atttacaaaa	240
atagtgcac	agcattatga	ataaactatg	aattggggac	catggaaatg	cactagaaca	300
aattttgtaa	aaatatggca	gatatggaag	ttaaaaatag	aatggatgca	aggactgtac	360
taaagggtgt	tggtgtagtt	acaatgttca	ctttgcacaa	ctatccctat	agtctaggta	420
qccattqqgt	ttctcctcag	cagtgtcaga				450

<213> Homo sapien

aaaaaatcat	aatgggggtt	tcataatcca	aagttgaaac	atttattctt	catagcttca	60
gaattttaaca	accaattgta	gaccatgctt	tccaaatcca	gtcttctttg	ctatttttca	120
aaacttctga	gatctagtat	taaactgctc	cattctaaat	gtatagtttt	agataagtat	180
tgtacacttg	ttgataaggg	ttttctgaaa	gcagtctatc	aatataaaag	aatgggttct	240
atctaagaat	cagcagtgag	ggaagaaata	ttaaacacct	atcaagaaat	caattattca	300
tttt						304

<213> Homo sapien

ctggaagaag	aactgagaca	gcagaaaagaa	gcagcttggt	tcaaggctcg	tccaaacacc	60
gtcatctctc	aggagccctt	tgttcccaag	aaagagaaga	aatcagttgc	tgagggcctt	120
tctggttctc	tagttcagga					140

<212> DNA

<210> 1397

<400> 1397

<210> 1398

<211> 261

<212> DNA

<213> Homo sapien

<400> 1398

aaaattataa	ctactcattc	tttcttttagc	cttagataat	ttgagcagaa	gccacaacaa	60
gcaaaccaca	ataaatttag	aattggcaga	aatccacatt	aactcctctt	cccaagtttc	120
cacactacta	ccatttacag	ttgtaggttt	gtaatgtata	attatgtaat	gcasaaacta	180
gctttgactt	gtgtracgat	gcactgtcaa	aggaagcaaa	gtaagaattg	aaattccaca	240
ttcccgaat	ttaacactca	g				261

<210> 1399

<211> 195

<212> DNA

<213> Homo sapien

<400> 1399

ctgatttttt	ttccttctca	aaaaaagtta	tttacagaag	gtatatatca	acaatctgac	60
aggcagtgaa	cttgacatga	ttagctggca	tgattttttc	ttttttttcc	cccaaacatt	120
gtttttgtgg	ccttgaattt	taagacaaat	attctacacg	gcattattgca	caggatggat	180
qqcaaaaaaa	agttt					195

<210> 1400

<211> 120

<212> DNA

<213> Homo sapien

<400> 1400

ctgcctccaa	ccctttgggt	ctccaccacc	caagtttcct	gtagggtccg	ccgggtccag	60
gatacacaggc	ctgggtttcg	tgagctgcct	tctcaggtag	ttttcaataa	tgggggttttt	120

<210> 1401

<211> 284

<212> DNA

<213> Homo sapien

<400> 1401
 ctgtagccaa aaagatgctg gggcagattg tggacaagta gaagcacctc cttccccctct 60
 gcgacattga acggcgtgga ttcaatagtg agcttggcag tgggtgggcgg gttccagaag 120
 gttagaagtg aggctgtgag caggagcctc tgccagggga catgcaatct gcagggaggg 180
 gctgaggggg gtcccatggt ctctgctgtc ttctctgtcc acctctttgt agaggagctt 240
 gagctccagg aatgctcttg tcagggctgc tgtgactggt ggcc 284

<210> 1402
 <211> 198
 <212> DNA
 <213> Homo sapien

<400> 1402
 ccaggtttct gctggtacca ggctaagtag ctggtgctgg cggaacact gtgactggcc 60
 ctgcaggaga ggggtggctct tcccccgga gacagagaca gcgtgtctgg agactgtgtc 120
 acttcaagct ctgcgatgcc atctgggagc cagagtagca ggaggaagag aagctgcgct 180
 ggggtttcca tggttccc 198

<210> 1403
 <211> 441
 <212> DNA
 <213> Homo sapien

<400> 1403
 aaactcaaaa ttgacaaatt aactagcttg ctttttgtca tttggaagac taccattatt 60
 caaattttatt atgtaataca ctcatccaga taatgaaaca tctgcgaaaa aaagtgtggg 120
 aatcacctca tctgtgcata aaatggctat tatacatgaa tgcagacgtt tgaagttaga 180
 aaggaatata actcaaatag caaaaggctc taattacaga gtttacaat aagcagtttt 240
 attttcaaaa gtacatagta agtccagact gggctattgc caaagaacta atcttttagtc 300
 tacttcaaca tgttacatgg tattcctgac tctacagact atcagcatct gtggaggtta 360
 gctcctaag gtcccaaaga acaggaaaca tgcaggaata aaggactcct catgaagagc 420
 aggtgggagc gagtgggcag g 441

<210> 1404
 <211> 243
 <212> DNA
 <213> Homo sapien

<400> 1404
 tgaaggggtt cttggaagac ctggcacctc cagagcgcag cagcctaatt caggattggg 60
 aaacatctgg gcttgtttac ctggactata ttagagtcac tgaaatgtc cgccatatac 120
 agcaggtgga ttgctcaggt aatgacctgg agcagttaca catcaaagtg acttcactgt 180
 gcagtcggat agagcagatt cagtgttaca gtgctaaaga tcgcctggct cagtcagaca 240
 tgg 243

<210> 1405
 <211> 168
 <212> DNA
 <213> Homo sapien

<400> 1405
 aaaccactgg atctatctaa atgccgattt gagttcgcga cactatgtac tgcgtttttc 60
 attcttgtat ttgactatct aatcctttct acttgtcgtc aaatataatt gtttttagtct 120
 tatggcatga tgatagcata tgtgttcagg tttatagctg ttgtgttt 168

<400> 1409
ccaqtccaac ctgctcctca ttattgtata aatgagcaga atcaatatgg cggaagccag 60

```

cttcaattgc caatttgggtg gcctctaaag ctttactttt aggaacctct gcaggcgcac 120
aggtgccaaa tcccaggaca ggcatgaagt gaccatcatt cagcttcaca cactgatatt 180
tcgaatccat ttctgtcact agcctggc 208

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<210> 1410
<211> 404
<212> DNA
<213> Homo sapien

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<400> 1410
aaaaaaagga aaaagtttta ttacgaaact agtttgtata aaacagggtt atacatattt 60
ttgtaagttt gtaataaaac agtaagaaaa aaaaggcagt aatagaaatc tccaaaaggc 120
aacctatcaa aaccaactgg ctgccacttt gagtttggac agtagctgca taaactttgt 180
tcttcttgar cagtatttaa taacatcatt aatacattaa caacatttct ataaagtaag 240
acacattggt gctgaagtac aactgggtgg ctcttgatct cacctatgag gagagttctt 300
tacamawcca catagggaaa attgcagttg taagggtgarc tacacatcta aaatatgcag 360
aggtaatagc attacatggt aaagtatcaa gatatacaca tttt 404

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<210> 1411
<211> 623
<212> DNA
<213> Homo sapien

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<220>
<221> misc_feature
<222> (1)...(623)
<223> n = A,T,C or G

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<400> 1411
ccacttggtg agatatgggg agcctacact ccggaggggst gtaccttttag cactggccct 60
catctctggt tcaaatccac gactcaacat cctggatacc ctaagcaaatt tctctcatga 120
tgctgatcca gaagtttcct ataactccat ttttgccatg ggcatgggtg gcagtggtag 180
caataatgcc cgtctggctg caatgctgcy ccagtttagct caatatcatg ccaaggaccc 240
aaacaacctc ttcattggtg gcttggcaca gggcctgaca catttaggga agggcaccct 300
taccctctgc ccctaccaca gcgaccggca gcttatgagc caggtggccg tggctggact 360
gctcactgtg cttgtctctt tcttggtatg tcgaaacatt attctaggca aatcacacta 420
tgtattgnat gggctgggtg ctgccatgca gcccgaatg ctggttacng tttgatgagg 480
agctgcggcc attgccagt tctgtccgtg tgggccaggc agtggatgtg gtgggcccagg 540
ctggcaagcc cgaaaactat cacagggttc cagacgcata caaccccagt gttggtgggc 600
ccacggggaa cgggcagaat tgg 623

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<210> 1412
<211> 171
<212> DNA
<213> Homo sapien

```

```

<400> 1412
gcggcgctgg ggggtgctgga gtccgacctg ccaagtgccg tgacacttct gaaaaatctc 60
caggagcaag tgatggctgt aactgcacaa gtgaaatcac tgacacaaaa agttcaagct 120
ggtgcctatc ctacagaaaa ggggtctcagc ttcttggaaag tgaaagacca g 171

```

```

<210> 1413
<211> 189
<212> DNA

```

<213> Homo sapien

<400> 1413

aaaagtcata	agggttttat	tttgtatcat	caaaatattc	tataagggtcc	caaatactct	60
ttttcaaccc	atgaacagta	agaatttgtg	aattctgata	atgaaaaaag	ttttcctcca	120
ggtatgtttg	tttcacattc	agtcctaaag	ccttgagcta	tgtgtacttc	cctcacacag	180
gaacaccag						189

<210> 1414

<211> 564

<212> DNA

<213> Homo sapien

<220>

<221> misc_feature

<222> (1)...(564)

<223> n = A,T,C or G

<400> 1414

cctccccagc	gccccaaagg	ctattacaag	tacctataga	cttttcacat	ataagttcta	60
gtgggtacaa	gctttttttt	tttttttttt	tttttttttt	tctattgggk	atttcattca	120
ttttgggggg	ggaacaaatt	ctacaaactg	ctttaatatt	gkcctttttt	tctaatactc	180
acattaactt	tttatgtaaa	acataccaat	gcttttaata	aagcttacat	aggaataaac	240
tattatagac	ctgcatagat	ataagtaccc	atgtattaat	ctacattaaa	ataatggatt	300
ttattctgcg	aaractccaa	gttgctcctg	ggkgctaagk	gaagcactta	gggaaatgtg	360
ttcagtcttt	gaggtcatag	gaacattara	ttatatcaaa	ggaaacctgg	agccatcagc	420
taagtggccc	ttctgtcctg	tagatacata	aaaactaatg	ggctccgcta	tgcggtcac	480
tttctgctat	tagatactat	gaggcactaa	naaaaaacta	ctgcttgcac	catatctttc	540
ttcggtttga	gataaagaga	atgg				564

<210> 1415

<211> 231

<212> DNA

<213> Homo sapien

<400> 1415

ctgcgcttgg	ataacaagta	attcaacgca	cgcacttaac	agaaatgtta	aactataaca	60
agcaccattt	gaggattaac	aggaacattt	ttttgaagat	ttcaaacgaa	ctcgactttc	120
agtataattg	tacctaaagt	atttataaac	agctcatcgg	agcctctatt	tgtcatagac	180
ttttgagttg	attgttggga	ccacataata	ggaccatttt	tttttgtctt	t	231

<210> 1416

<211> 540

<212> DNA

<213> Homo sapien

<400> 1416

cttgatttag	gatctgtggt	gcagggcaat	gtttcaaagt	ttagtcacag	cttaaaaaaca	60
ttcagtgtga	ctttaatatt	ataaaatgat	ttcccatgcc	ataattyttc	tgtctattaa	120
atgggacaag	tgtaaagcat	gcaaaaagta	gagatctgtt	atataacatt	tgttttgtga	180
tttgaactcc	taggaaaaat	atgatttcat	aaatgtaaaa	tgcacagaaa	tgcattgcaat	240
acttataaga	cttaaaaaat	gtgtttacag	atgggtttatt	tgtgcatatt	tttactactg	300
cttttcttaa	atgcatactg	tatataattc	tgtgtattttg	ataaatattt	cttctacat	360
tatattttta	gaatatttca	gaaatatata	tttatgtctt	tatattgtaa	taaatatgta	420

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catatctagg tatatgcttt ctctctgctg tgaaattatt tttagaatta taaattcaca      480
tgtcttgctca gatttcatct gtataccttc aaattctctg aaagtaaaaa taaaagtttt      540

```

<210> 1417

<211> 350

<212> DNA

<213> Homo sapien

<220>

<221> misc_feature

<222> (1)...(350)

<223> n = A,T,C or G

<400> 1417

```

ttnatcatct aactgtggga tctatttcat ttctggaaat aacacaactt agttctaggg      60
ctttcatgca catgaaatat aaaacagctt agttgttctg aaaacatgac aatggttaat      120
tttattcaag tccaacact gagttcagag cacttctcca taggccccat taatctctcc      180
aggtttctgg gagtatcatt aaatccctcg gcattcctta gaagcaggtg cttagcaaac      240
atccagtttc caaatgagag tcagaggggc ttgatcctga aagtgtagta ttttcttgcc      300
ttgtctact ggtatagctt cttggaccta aaatctctct cctgctgagg      350

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<210> 1418

<211> 425

<212> DNA

<213> Homo sapien

<400> 1418

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tgctaggcag ccttattttc ataaccawt tagggaaagg aaatttagga ttttcaaggc      60
tacattaatt tttctccat caaatcttga ttgttcttg ataaaaatga gttcttttgg      120
ggaaattctt tctttagaca ccaacttggg ttttctcatc tccacagaa taattgaacc      180
cctgacctct agatgttcaa aattccgctt caagcctctg tcagataaaa ttcaacagca      240
gcgattacta gacattgcca agaaggaaaa tgtcaaaaatt agtgatgagg gaatagctta      300
tcttggttaa gtgtcagaag gagacttaag aaaagccatt acatttcttc aaagcgctac      360
tcgattaaca ggtggaaagg agatcacaga gaaagtgatt acagacattg ccggggtaat      420
accag                                           425

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<210> 1419

<211> 390

<212> DNA

<213> Homo sapien

<400> 1419

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aaactcttgc tattgaattg agatgattaa aatggtgact taatccgtag ttattttgca      60
cccactgaaa ggaaagtgtt ttccagaata atatgaagta tctaaaagtg tcaccttttc      120
ttgcctgata aacaatttgg gcttctgtt tgtacaaggg gccatttggc atacctttca      180
cagcttttat caggccaagt taaaggctga ctacattttt tcatcatgag gaaagcagtt      240
gaaatgaggc atgagttact gtgcattggg atttttagaac aattttcttg tgacagctct      300
tttgtgaag ttaggttctt aaaagtgcc atgatggtca cttaaaatgt gcagtaatag      360
cactgccagg atcaagcatg aaaggctttt                                           390

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<210> 1420

<211> 480

<212> DNA

<213> Homo sapien

<220>
 <221> misc_feature
 <222> (1)...(480)
 <223> n = A,T,C or G

<400> 1420
 ttgctgaaca atgacatcgt tttctccagg ggttgaaatc catgtccatg gctgacaacc 60
 caacaaggct gggacccaaa ttcgtacaga gatgaggcag agtggagaga aacaactctg 120
 gctgagccag agtctccagc cactacttct tattcctggg ctttagctct tcggctgcat 180
 tacgcaggaa aatgtaattt tttttctggg gattataaaa ttcattgtccc tttgaccagt 240
 cgtagctgga agcgtatgca aatatgtttc cattgygatt gaaacagcaa gctgasatgg 300
 gctgayctaa ctgttccgaa gnttttagtt ttgktctggc atctttgycc cagaagctga 360
 atctaccatc agatcccaca gttgcaaggg tgccatgaac aggatggaac gccgattcca 420
 tttaccgcga taaatgycct gaggagctga agtggttggtt ccattagatc gatgacattt 480

<210> 1421
 <211> 453
 <212> DNA
 <213> Homo sapien

<400> 1421
 aaactgattg aggtcacagt attttattat ttggggtoct caccacagga aacactgcga 60
 tacaggggca aaagagatgg cagtgccaat taaattaata caacaaaatc aatgcagcac 120
 caaccaagac tgccaggtct ggtgtcatgg gtatgccag agcccaggag ttcagaaggg 180
 ccctaagcct gatttaatgc tctgctgttg atgtcttgaa attcttaaca atttttgaac 240
 aaggggcctg cgttttcact tcgcactggg ccttgcaaat tacatagcga gtgctcataa 300
 aagaactcag aaacgtggta cctctcttcc tgggtggatac aaataaagaa atctggatcc 360
 aaagttgaaa gttgctggcg atatcattca agtaggactc taaatagtgg attaagatga 420
 ggggtgggcct ggggtgaagat tctttccagc ttt 453

<210> 1422
 <211> 542
 <212> DNA
 <213> Homo sapien

<220>
 <221> misc_feature
 <222> (1)...(542)
 <223> n = A,T,C or G

<400> 1422
 ttttcttgac cactatacgg cacaacctag gggstgtawa aaacctascr caatgcagaa 60
 ggggtgaagct tcatgacaat tggctcgcgc aataatttgg gggatgtaac atcaacgaat 120
 cagacaacaa aagcaaggga atacacatgg nactaaatca gtgtgnggaa aaatatccca 180
 aacaggcaaa gcacaacatg gamtagatat atgcacattn atggaccctg naggcakkac 240
 tcacaaacat actacctggg aagcamctgg acctttaagg gatgaggtag attcaacaaa 300
 cagggcancg tatmttcac tgggatagca ttccagcctt aaaaataang aaatcttgaa 360
 aagnactaca ataaggacaa atctcgaaca cattctgtta agtaaaacaa gacaagccaa 420
 aaagggaana ctgtataatt acacctatgt aaaatattta gtcaaaactca aagaaaccaa 480
 gtgttgtagt ctgagcaggg caccaagatg naaacagtct ctcatagnct gagatangca 540
 tc 542

<210> 1423

<213> Homo sapien

ttaatgccaa	atggccaaagt	tgcattccgtg	gaaatgggta	aatatcatca	ctgtcgggat	60
gaaccctctg	acgccctcta	tgacaatgtg	gagaaactct	ttccagggtt	tgagatagaa	120
actgtgaaga	acaacctcag	gattcctttt	aataatgctg	taaagaaacg	tttgatgaca	180
gacagaagga	ttggctgcct	tttatcaggg	ggcttgga	ccagcttggt	tgctgccact	240
ctgttgaagc	ag					252

<213> Homo sapien

tttccactct	gcacattgta	gagggaaacac	tctgtaggcc	catgggtccc	ttactagaga	60
ggttgagtga	atttgccctc	agttaacatg	ggaccttctg	ttagcttcc	tcttgcttcc	120
caaagatttt	aagcattttg	taaatgtata	aactcacctc	tggtaacagt	ggcccagacg	180
ctgctttgtg	ctaaaagcat	gggaatgta	aaggcagctc	ttctctggga	aatggatgct	240
attctattct	gtagccccta	gctgttctcg	agg			273

<213> Homo sapien

aaaaaccttg	tatagcaaaa	taacttaaaa	ccctttgtga	tatcatctta	ccagttttatt	60
tggtaaaaaac	aaacagttat	ttggtatttg	tcagaattct	tcagtgcctg	ctattacagc	120
tattttccaa	ttactaattt	gattatactc	actcaaggca	gtgcaagatc	ttgaagtact	180
tttttagcagt	taagtaatat	tgaattgtat	tgaatagttt	acatagttta	ttctagtctt	240
tgaaaattac	tgaacatgga	caatgtgcat	gtcattgaca	tctgccttag	aacttctggg	300
acaatcctga	ttcgagagat	tctatcccat	tatttacata	tacaaaaaat	actttgttaa	360
tttaatgtgt	tggtcttcca	actcctgaac	acgacacaaat	tttattatta	gattttgtat	420
ggtgattttt	ggctatgaaa	acatgatcat	tatatgata	tagatacatt	tttatttgtt	480
acaaatgttt	gagcagctca	ctagcccacc	cctcctctat	tttgggtaag	agaatttact	540
acctttttta	actatgtagt	tgagagcaac	atgtattttt	ttatttttag	aatggtcagt	600
atattgctat	aaaatttt					618

<213> Homo sapien

gtggtagaaa	gagatgacgg	aagcacatta	atggaaatag	atggcgataa	aggcaaacaa	60
ggcgggtccca	cctactacat	agatactaata	gctctgcgtg	ttccgagggga	gaatatggag	120
gccatttcac	ctctaaaaaa	tgggatggtt	gaagactggg	atagtgttcca	agctattttg	180
gatcatacct	acaaaatgca	tgtcaaatca	gaagccagtc	tccatcctgt	tctcatgtca	240
gaggcacccgt	ggaatactag	agcaaagaga	gagaaactga	cagagttaat	gtttgaacac	300
tacaacatcc	ctgccttctt	cctttgcaaa	actgcagttt	tgacagcatt	tgctaattggt	360
cgttctactg	ggctgatttt	ggacagtggg	gccactcata	ccactgcaat	tccagtccac	420

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<210> 1427
<211> 144
<212> DNA
<213> Homo sapien
```

```
<210> 1428
<211> 214
<212> DNA
<213> Homo sapien
```

```
<210> 1429
<211> 253
<212> DNA
<213> Homo sapien
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<220>
<221> misc_feature
<222> (1)...(253)
<223> n = A,T,C or G
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<210> 1430
<211> 232
<212> DNA
<213> Homo sapien
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<210> 1431

<211> 734
 <212> DNA
 <213> Homo sapien

<400> 1431
 cattatacaaa cactatatattg ccagggtcaaa gagggcaggg acgtaaatgt acactaaaat 60
 gcmaatgtat cccaaagaga taaaacaaat tccattttaca gcatgaaggt ttacaaatgt 120
 acacctgtac aaccaaggaa agcatcacta ctaaattagc aaggctttta taataaacat 180
 tgaaasaaga tttcctttca aagtgtaaac ttacatctat tactacacac acaatgcata 240
 tattttataga aagcaaaaag agctatctga atatgtaatc atgcttaaat gctgagctat 300
 caaattcact tttcagtggc cccttttcat ctctatctgg ttctactttt ctgcctctat 360
 gaaaaagcaa aataaagctc aacacttcct caacatgtct gtaattctat aagcaaaaca 420
 aaatacaaat ttccactctt tctcattgca aaccaaactg aaaagttaat aagtgactta 480
 acttttcatt tagtgacttt aattggaagt gtcaccatga ttttgtattt aactcttaca 540
 acaattacat atgttaagtat atacaatatt tctgtacatt gccagagaca ttttagggca 600
 gtaattgtat taaaaccaca tctactgtaa ataatgttag gttctttttca tctcaaacca 660
 ctttattctt gcctacttac tcgttatattg catgatagtt tgtgaattat caaaatacaa 720
 cttaactctt taaa 734

<210> 1432
 <211> 542
 <212> DNA
 <213> Homo sapien

<400> 1432
 tttaagaaag agccttttgag aaacatgcat actttttctct tttctcctat attcaatact 60
 catatagcct aaaagatgga aactgggttca agaattttaa tgacttggtc cctaaaaagt 120
 taatctcttc accctttgtga aatatatcaa gtgcttttcta taaataaggg caggaaatgc 180
 taacttcata agcatagtcc tagtcattaa aataatttga tcatcttcta aattttaagt 240
 atgatagtaa cacagtaata tggaaaatct caatatactt aacacttcct aaacagcaca 300
 atgaaatggt gttcaaggtc tgaattaatt tgctacagga cctaagcaag tctgtttgct 360
 tatcttttgg ctttaaaatt ctttaagtct aaaatgggtga taattttaga ataaactgac 420
 aatgtgggga acaaacttaa attcacaac actaccata tgctcaaaaa ctctctggga 480
 taattagttt cttcattgta actattgatg tactattatt tcatctttcc attagctcta 540
 ct 542

<210> 1433
 <211> 175
 <212> DNA
 <213> Homo sapien

<400> 1433
 ttaaattgat tcaaaaaaac ttgacacctg tcatgtaggc caaaaaatag tagcgaacta 60
 tactaagtgg tatagcccac tgtggagtgt ggtcttttac tcttccaaat agcccaagtt 120
 ggcaaagggt acttaaaaaac ctgccccca aaaagctaac ttttggtaga ttttt 175

<210> 1434
 <211> 90
 <212> DNA
 <213> Homo sapien

<400> 1434
 ttaatcacta ttgatggaag cttatatctc ttatgaatat atacatgtat gcatatatac 60
 atctctgtat gaatcactca aagcaatttt 90

<210> 1435
 <211> 153
 <212> DNA
 <213> Homo sapien

<400> 1435
 tttacctttg tgctttgaag gttctaccat ttakaaagta aaaagccaac ccacagaatg 60
 gaagaaaaga ggacagactc taacaagcgt tcacaaagat ggagagaaat tgtaaccctc 120
 atatattgct ggtagaattg tagaaagatg cag 153

<210> 1436
 <211> 483
 <212> DNA
 <213> Homo sapien

<220>
 <221> misc_feature
 <222> (1)...(483)
 <223> n = A,T,C or G

<400> 1436
 ttttttagttt aaagaagagt tttgccactt aracanggga gctwtgtctg gaaaatacac 60
 tgagttgaaa cacttcaccc ttggaaggat tatataagat gaacagytgt gataaatgtg 120
 tagattagag ggatgtgaat gggcagttag tccagtgcc tcatttaaga ggccaagatc 180
 ctgattcaga ggaggcatcc tttgccaga gctgcttagc taatctgacc aaatgttggg 240
 aaaaatgtct cacctaacc actattcctt aattatggat tttgtgaaaa acaatagaac 300
 atgttaatga gtaatttata ttagttcgat gtattacaat tttttagctt taaattacag 360
 ytttcttata atgttgaaat gttttagaat cctttgaatc taagtatttg tttcctaaat 420
 gaaacatttg tacaacattt gatgttttta cttatgaaat attctcctcc cccaagaaaa 480
 ttt 483

<210> 1437
 <211> 171
 <212> DNA
 <213> Homo sapien

<400> 1437
 ttttgccacc tcaagaagcc attttcttgt ctgtttcctt ctttacctac ccctacaacc 60
 tatgaacaaa taccataact taaaaattta ggtagtctac aactcctaca aatttttaagt 120
 tcagagacta cccaagaac tgtggaagat gcagcaatat aaaagttttt t 171

<210> 1438
 <211> 408
 <212> DNA
 <213> Homo sapien

<400> 1438
 tctgagtgga ggtaggctaa caacacattt tgactttstc ctcaaaggat agctttgaaa 60
 aacaagtgta accaattgtt acaccaaat aaaatggcaa tattaatcg gtaacaaaac 120
 gatccacatt ttatacaata ttgtatttcc aaacatacat aggtcatgaa aatcagagaa 180
 cctaatatag caccgttgaa accattcatt atccttcatt tgtgtatgca attcagaatt 240
 tcggcagaag acaacaaatg gaaaatgcct ttcgtttcta taaatcattt tggatttcaa 300
 ttaaattctt gccttagtaa agggatttct tatctcaaga tcaattagcc gtttttagct 360

ccaccgtttt ggaagtaaaa atgatgagct acatctactt ttttaattt

408

<210> 1439

<211> 168

<212> DNA

<213> Homo sapien

<400> 1439

ttacacaaca	gctataaacc	tgaacacata	tgctatcatc	atgccataag	actaaaacaa	60
ttatatattag	cgacaagtag	aaaggattaa	atagtc aaat	acaagaatga	aaaacgcagt	120
acatagtgtc	gcgaactcaa	atcggcattt	agatagatcc	agtgggtt		168

<210> 1440

<211> 307

<212> DNA

<213> Homo sapien

<400> 1440

tttcacatac	gaagaaatca	actgtgatta	tgaagtgaca	gccagctaaa	tatgtcttgt	60
atcttctctc	ttcctttttt	tgctaaactc	atcctttact	tccattcctg	cttccatggg	120
aatgcaggct	caaataaatt	actaggatac	aagattactt	caagcctctt	ttctgtggaa	180
ctcataatat	gataagcatt	tgttacaaga	ttgcctgtag	ttgtttaggg	gacaaattat	240
attagggaaa	gaaagtcttt	cttttagttg	ttaaattttc	tattataatt	gggtactaaa	300
tttattt						307

<210> 1441

<211> 684

<212> DNA

<213> Homo sapien

<220>

<221> misc_feature

<222> (1)...(684)

<223> n = A,T,C or G

<400> 1441

ttaagttctg	gagtgttcac	ttctgagcct	gaattccctc	ccctgcaaaa	tgggggaata	60
ccctcctcag	aggggtccctg	cgagggtgag	gggagattca	gcatggcagg	tgtgctgggc	120
acggcagggc	ctgggaagg	cagatccttt	ccccatccct	gccacaaaca	acccaaacct	180
ttaaaggaga	gcaatggcct	tgtgtcaaaa	acaaaaacaa	aacaaaaccc	tgtcctagga	240
gactggggcc	ctaatttcta	atagcaagcc	tttatgagtc	cctaacactc	tactgggctg	300
agtatctcac	acgccagagg	ataacctgcc	ttctgctcac	caccaccccg	tagtagttgt	360
cattgtgtcc	atttcacaga	tgaggcaaag	gctcagaaga	gtcatgtgtt	aaaccagctt	420
ctagagccca	tgcaggagct	gcagggtggga	gaatcacctc	taggtgctct	tcccatagaa	480
tcctcacctc	ctgagtgtca	ctcactcagc	ttccaatggg	tgtgtgacct	ttgaccagct	540
ttcttctctc	ctgggcctca	gtttccacc	tggacaaagt	aagaggtctc	ttggcttcan	600
gtaagttctt	cctaaacttc	tttttccttt	tcatttgagc	atcctcttca	tttttgccac	660
ctctctgtca	tttacaggct	tttt				684

<210> 1442

<211> 166

<212> DNA

<213> Homo sapien

<400> 1442
 aaaaaatcag cccctaattt ctccatgttt acacttcaat ctgcaggctt cttaaagtga 60
 cagtatccct taacctgcca ccagtgtcca ccctccggcc cccgtcttgt aaaaagggga 120
 ggagaattag ccaaactg taagctttta agaagaacaa agtttt 166

<210> 1443
 <211> 194
 <212> DNA
 <213> Homo sapien

<400> 1443
 tttgccctgt caaaagaaga gctaaagaca gttatataaa aattaagggtg ggctttcaga 60
 ctggctaaca caacaacatt ccatgagtag atggtaattt atttttgttt atccatttcg 120
 ttgggagcaa ggacaaaaat gtaaactctac accttgctta tcaaaattgc cgaaaaaaga 180
 atgctctgcc tttt 194

<210> 1444
 <211> 96
 <212> DNA
 <213> Homo sapien

<400> 1444
 gagagtgcag agtgggagaa gagcggagcg tgtgagcagt actgcggcct cctctcctct 60
 cctaacctcg ctctcgcggc ctacctttac ccgccc 96

<210> 1445
 <211> 365
 <212> DNA
 <213> Homo sapien

<400> 1445
 gggatgagct gaccaagaac caggtcagcc tgacctgcct ggtcaaaggc ttctatccca 60
 gcgacatcgc cgtggagtgg gagagcaatg ggcagccgga gaacaactac aagaccacgc 120
 ctcccgctgt ggactccgac ggctccttct tctctacag caagctcacc gtggacagga 180
 gcaggtggca gcaggggaac gtcttctcat gctccgtgat gcatgagggg ctgcacaacc 240
 actacacgca gaagagctc tccctgtctc cgggtaaatg agtgcgacgg ccggcaagcc 300
 cccgtcccc gggtctctgc ggtcgcacga ggatgcttgg cacgtacccc gtgtacatac 360
 ttccc 365

<210> 1446
 <211> 386
 <212> DNA
 <213> Homo sapien

<400> 1446
 tctggaaagt tcttgctcgg gtcccttcac ctccccgcc tttcttarag tgcagttctt 60
 agccctctag aaacgagttg gtgtctttcg tctcagtagc cccacaccca ataagctgta 120
 gacattgggt tacagtgaac ctatgctatt ctacagcctt tgaaactctg cttctcctcc 180
 agggcccgat tcccaaacc catggcttcc ctacactgt cttttctacc attttcatta 240
 tagaatgctt ccaatctttt gtgaattttt tattataaaa aatctatttg tatctatcct 300
 aaccagttcg gggatatatt aagatatttt tgtacataag agagaaagag agagaaaaat 360
 ttatagaagt tttgtacaaa tggttt 386

<210> 1447

<211> 261
 <212> DNA
 <213> Homo sapien

<400> 1447
 aaaattataa ctactcattc tttcttttagc cttagttaat ttgagcagaa gccacaacaa 60
 gcaaaccaca ataaatttag aattggcaga aatccacatt aactcctctt cccaagtttc 120
 cactacta ccatttacag ttgtaggttt gtaatgtata attatgtaat gcagaaacta 180
 gctttgactt gtgtaacgat gcactgtcaa agtaagcaaa gtaagaattg aaattccaca 240
 ttcccagaat ttaacactca g 261

<210> 1448
 <211> 404
 <212> DNA
 <213> Homo sapien

<400> 1448
 aaaaaaagga aaaagtttta ttacgaaact agtttgtata aaacaggggtt atacatattt 60
 ttgtaagttt gtaataaaac agtaagaaaa aaaaggcagt aatagaaatc tccaaaaggc 120
 aacctatcaa aaccaactgg ctgccacttt gagtttggac agtagctgca taaactttgt 180
 tcttcttgaa cagtatttaa taacatcatt aatacattaa caacatttct ataaagtaag 240
 acacattggg gctgaagtac aactgggtggc ctcttgatct cacctatgag gagagttctt 300
 taaaaacca catagggaaa attgcagttg taagggtgaac tacacatcta aaatatgcag 360
 aggtaatagc attacatgtt aaagtatcaa gatatacaca tttt 404

<210> 1449
 <211> 230
 <212> DNA
 <213> Homo sapien

<400> 1449
 aaaagttcta gtggtacggg aggagctttg caggaagttt gcaaaagtct ttaccaataa 60
 tatttagagc tagtctccaa gcgacgaaaa aaatgtttta atatttgcaa gcaacttttg 120
 tacagtattt atcgagataa acatggcaat caaaatgtcc attgtttata agctgagaat 180
 ttgccaatat ttttcaagga gargcttctt gctgaatttt gattctgcag 230

<210> 1450
 <211> 194
 <212> DNA
 <213> Homo sapien

<400> 1450
 aaaaactcct tttggtttac ctggggatcc aattgatgta tatgtttata tactgggttc 60
 ttgttttata tacctggctt ttactttatt aatatgagtt actgaagggtg atggaggtat 120
 ttgaaaattt tacttccata ggacatactg catgtaagcc aagtcatgga gaatctgctg 180
 catagctcta tttt 194

<210> 1451
 <211> 106
 <212> DNA
 <213> Homo sapien

<400> 1451
 aaagatgaca aatactggtt aattagcaat ttaagaccag agccaaatta tcccaagagc 60

atacattctt ttggttttcc taactttgtg aaaaaattg atgcag

106

<210> 1452

<211> 349

<212> DNA

<213> Homo sapien

<400> 1452

ctgcagatcc	tgcggaacgt	caccaccac	gtttccgtga	ccaagcagct	cccaacctca	60
gaagccgtgg	tgtctgctgt	gagcgaggcg	ggggcgctctg	gaataacaga	ggcgcaagca	120
cgtgccatcg	tgaacagcgc	cttgaagctg	tattoccaaag	ataagaccgg	gatggaggac	180
tttgctctgg	aatctgggtg	tggcagcatc	ttgagtactc	gctgttctga	aacttacgaa	240
accaaaacgg	cgctgatgag	tctgtttggg	atcccgctgt	ggtaacttctc	gcagtcctccg	300
cgcgtggtca	tccagcctga	catttacccc	ggtaactgct	gggcattta		349

<210> 1453

<211> 302

<212> DNA

<213> Homo sapien

<400> 1453

aaaaataatg	tgcaagagca	tcatgagaaa	gaagaggggt	gaagagataa	tccagaggaa	60
catcaaagt	aagagtatac	actcaaagac	aggtttaaga	aagaccagtc	agagaagtaa	120
agaaaaaat	caagcaagaa	taatgttgca	aaaattaaca	agaaagttgc	aagcccagag	180
tggttagcaa	tgccaaacta	ccatgagtaa	gccacataaa	acaagaactt	tgggttcaac	240
tgctttaaca	atcagacctt	tagattcaca	taacaggagt	tacaaaatta	agagcctctt	300
tt						302

<210> 1454

<211> 268

<212> DNA

<213> Homo sapien

<400> 1454

caagcgtaaa	ccgcgggagc	cgagcccagc	taggaatgca	gacctcctga	aaaccaagcc	60
gaggactgcg	gggtccgggtg	tccacgcaga	gtgtcagctt	cctctggtgc	aaccagcaag	120
tcttccagta	tgaatcccac	agaaaccaag	gctgtaaaaa	cagaacctga	gaagaagtca	180
cagtcaacca	agccaaaaag	cctacccaag	caggcatcag	atacaggaag	taacgatgct	240
cacaataaaa	aagcagtttc	cagatcag				268

<210> 1455

<211> 207

<212> DNA

<213> Homo sapien

<220>

<221> misc_feature

<222> (1)...(207)

<223> n = A,T,C or G

<400> 1455

ctgtcgagag	cagccctgcc	caagawtgnc	gggtgggggc	tggtgccaac	gggttcccaa	60
ggscctttcm	actttkgaak	ggctggartt	cttgggaaac	cmaaacsctg	actacctgsc	120
ttttttcttg	ggcatygacs	tgcttcattt	ccaaaratga	tggkgcaggt	gaccttttcc	180

atcgtgagct aaaaaaaggt taggagg 207

<210> 1456

<211> 181

<212> DNA

<213> Homo sapien

<400> 1456

aaatttctgt	ctgctaaaaat	ctatcaaata	cattaaggaa	aagtcccact	tggcacatct	60
cccacaccag	atgttaatta	ttcatactgc	atgactgagg	attttggagg	cagagagaga	120
ttcatctgca	atatttggaa	caccaatgga	ggtctacgtc	aacacagaat	ttatacagca	180
g						181

<210> 1457

<211> 309

<212> DNA

<213> Homo sapien

<400> 1457

aaaaagwtca	gagttgaaat	gcctttcaac	cattkecttc	tgtggtcatt	tttcttgctg	60
cctttttcac	ccaagattca	gcagtcagat	gtttactgca	cacctattac	ctattatttg	120
ctgttcttgc	atgggtcaaa	ccaccattct	gtagccaccc	atcctttgcc	ttatctaaca	180
aacatttttc	caggaagggtg	gaaaaggaag	tgttgctctc	attgtgtgac	tcagtgtctg	240
tgtccatccc	atggaaacat	gggcacaatc	aagtattttgt	ccagcctatt	gcaggctttt	300
cctgacttt						309

<210> 1458

<211> 117

<212> DNA

<213> Homo sapien

<400> 1458

aaagactatt	gagaaatagg	aaggtattga	gagattattg	ggtttcatca	kagcagactt	60
aagtagcctg	gttgatttta	gatttgtcac	agcaaaatca	tgcttggatg	ctcgagg	117

<210> 1459

<211> 575

<212> DNA

<213> Homo sapien

<220>

<221> misc_feature

<222> (1)...(575)

<223> n = A,T,C or G

<400> 1459

aaagaatgca	taccagaaca	tttataagca	gtggagtgag	kthtattaag	aatagtacta	60
ctacaataaa	cgctggctaa	ataagaagt	cattatgtga	agcactatgg	gtggatatatg	120
cttwgmcaca	tactctkgtt	accttgaggy	agatmacrca	tgkgaaccaa	cttcggcata	180
catttttcagt	tgtctgcgagg	aatcatgtgt	tttaacgaaa	tgctgcagta	tgaaaaactt	240
gaaaatattc	atgaatgawg	aacgcmntag	gaaaaaaata	kstattctca	tgcaattatg	300
tacagtctca	ctgtgtarat	ctcaaggcaa	ggtttgctc	ctgtaaacca	gatcaagggtg	360
ctatgagaga	ncgccytgnc	ttattgcatt	tcttttctcc	tmctgcgcca	gcattatatt	420
gctctagnct	ttatttttgt	gtgcacactg	acatgccatt	aaaratgang	ractatctca	480

catgtagaaa argaaagnmc ttgganketa cctcaggtcg ctaccacgct aaggggyaat 540
tctgcaggat atccatcaca ctggcggcgc gattg 575

<210> 1460

<211> 444

<212> DNA

<213> Homo sapien

<400> 1460

ctgggggttc	cttccttcac	gttgagaacc	tggagcagag	agtctacca	cttaagaaat	60
attagaaaga	gttcagcaaa	cagagtgcgc	tgaagtctaa	tcctagaagt	aaatccattc	120
ctacaagtca	tcagcatcac	ttgggagctt	gttagaaagg	caaattcttg	gttcagccta	180
acacctaata	aatcagaaac	tctgggggag	gagcgagca	atctgtactt	tcacaagccc	240
tgcaggtgat	tctgagcctg	taaaatttga	gaaccagagc	tgtccccag	gagataaatt	300
aacttctact	tttttttgag	ctactgcatt	ttgggatctt	attgttttat	cagcttaaca	360
tgcactctga	tatgattact	caggtatggt	tcaaccaatg	ttgggttaatg	tattatcccc	420
aggaacttat	tactagagga	gcag				444

<210> 1461

<211> 536

<212> DNA

<213> Homo sapien

<400> 1461

ctgcaaccct	gggactgacc	gggaggctct	gattatttac	ccmaccacag	gtaggttggtg	60
ttctgaatct	caggttcaca	ggttaagggt	cagcatcctc	atcctccacg	gggttgaggt	120
tgttgctggt	gatgaagggt	ttgggtggct	ctgcatagac	tgtgatcgtc	gtgactgtgg	180
tcctattgag	gccactggct	gagttattgg	cctggcaggt	atagagtccg	ctgttcttct	240
cagtgatgtt	ggagataaag	agctcttggt	tgtgttgctg	gatgttccca	tcaatcagcc	300
aagaataact	tgcaggtggg	ttagaggctg	catggcagga	gaggctgagg	ttcaccctg	360
gacggtaata	ggtgtatgag	ggggaaatgg	tggggkctc	ygggccatag	aggacattca	420
ggatgactgr	gtcgctgtgs	tyarcactta	atkcgttctg	gattccacac	tcatagggtc	480
ctacatcatt	ccttgtgaca	ytgartagag	tgagggtcct	gttgtcattg	gacagm	536

<210> 1462

<211> 409

<212> DNA

<213> Homo sapien

<400> 1462

ctgakagacc	aggagaagtt	ccagatgcag	agactgtgat	gctcttgact	atggaattat	60
tgcggccagt	agccaagtta	gagacaaaac	aggcataggt	cccgttatta	tttggcgtga	120
ttttggcgat	aaagagaact	tgtgtgtgtt	gctgcggtat	cccattgata	cgccaagaat	180
actgcgggga	tgggttagag	gccgagtggc	aggagaggtt	gaggttcgct	cccgaagggt	240
aagacgagtc	tgggggggaa	atgatggggg	tgtccggccc	atagaggaca	tccagggtga	300
ctgggtcact	gcggtttgca	ctcactgagt	tctggattcc	acatacatag	gctcttgcgt	360
catttcttgt	gacattgaat	agagtgaggg	tcctgttgcc	attggacag		409

<210> 1463

<211> 502

<212> DNA

<213> Homo sapien

<400> 1463

ccttcagcct	ggatccttta	tattaagatc	aatgaggacc	atttctggaa	gatgtctggc	60
atggtacaga	ctgtctgagg	ccractgaac	acaggccctt	accctgattt	tatcagtga	120
aagctatggg	actagtttcc	ttacctctaa	aatggagaga	ataatagaat	cttccgtcta	180
agactkctgt	gagcataagc	cgagaaaatg	gaggtaaact	gcttagccca	atacttggat	240
tatcgtaa	attcagtaaa	actagccacc	gttggtattg	taattattat	tttgtatttt	300
attatacatt	tcattgaaa	ttaaaagtta	gtgataatca	cctcattttc	agttgccttg	360
ctttcttct	gtaaatttta	ttctctctta	tcttgetcac	tgtctttaag	cattgccagt	420
ttagtataat	tattttcccc	tatcctctat	aaaatcatat	acaggatgga	tttgttgatc	480
tcagacatgt	tcactgagtt	tt				502

<210> 1464

<211> 294

<212> DNA

<213> Homo sapien

<400> 1464

ggcggctcgg	actgagcagg	acttttcctta	tcccagttga	ttgtgcagaa	tacactgcct	60
gtcgtctgtc	ttctattcac	catggcttct	tctgatatcc	aggtgaaaga	actggagaag	120
cgtgcctcag	gccaggcttt	tgagctgatt	ctcagccctc	gggtcaaaaga	atctgttcca	180
gaattcccc	tttccctcc	aaagaagaag	gatctttccc	tggaggaaat	tcagaagaaa	240
ttagaagctg	cagaagaaag	acgcaagtcc	catgaagctg	aggtcttgaa	gcag	294

<210> 1465

<211> 249

<212> DNA

<213> Homo sapien

<400> 1465

gtgcaggtct	tcagccgtga	cccgttacc	cagctctaag	ggaggtggca	gcatcaaagg	60
ctccccctgc	ctgcgtggca	gcagggaat	cttgctgcta	cggggcctag	agtcattggga	120
tctgggggag	ccaccctgg	gggcaagtgt	ctgccctggg	gctgtacctg	ccttgttttc	180
acagcgggtga	cccgaagaga	cagcctgagg	tccgtcctca	ctcactgtgt	ttgaggaact	240
gtggggccag						249

<210> 1466

<211> 203

<212> DNA

<213> Homo sapien

<400> 1466

cctcagacac	cttttaattg	cttaggagaa	accattgtct	ctgactgcag	gtttgaataa	60
gttgaagacc	agagaaaagt	acacactggg	ctacaaagga	atttgagat	agccaaggaa	120
caggatttcc	cctagcaagc	tacettctgt	tcaaatcatg	aaaaaagact	atttccccct	180
agaataggga	agcttgctat	ttt				203

<210> 1467

<211> 223

<212> DNA

<213> Homo sapien

<400> 1467

ctgtcagaac	aggaacgacc	tgggttatgg	aagcccagaa	agggaggagg	acttcttttg	60
gtcccagtga	aagatgcttc	cagaatctgt	agccttactt	atttgcttgg	atctcactgg	120
aataacttgg	tggtgaggtc	accggttctg	gggtgatcac	tgggtttgct	gcatagatgt	180

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<210> 1468
<211> 177
<212> DNA
<213> Homo sapien
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<210> 1469
<211> 185
<212> DNA
<213> Homo sapien
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<210> 1470
<211> 482
<212> DNA
<213> Homo sapien
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<210> 1471
<211> 257
<212> DNA
<213> Homo sapien
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<210> 1472
<211> 342
<212> DNA
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<213> Homo sapien

<400> 1472

cttttgcgag	cctctgccgc	agcagctccg	ttttcacgcg	catctcgttt	ttgtgtgtgt	60
gtttttgttt	tgtttttgtt	tttgtttttt	tgtttcagag	aattggaagc	ttaaagctacc	120
aaagacgtag	aaagaaatct	tagcaggtaa	gatggggcgag	ctttccgtct	cccggcccac	180
gataatcgta	tattttctact	ccgattcgcc	ctttctgggt	tgagaagttc	ccccgtgaca	240
ttttcttccg	cacccggaga	gcagacattc	gggagaagcg	gcctggggga	atactggagg	300
gattgcgggg	agatgcgtaa	ttacgcgtgt	gtttctttct	tt		342

<210> 1473

<211> 526

<212> DNA

<213> Homo sapien

<220>

<221> misc_feature

<222> (1)...(526)

<223> n = A,T,C or G

<400> 1473

ctgctacatg	tcttcacagc	ccaggaattc	aaggcccagg	tggcagcagg	aagaaacagt	60
ggaaaagcaa	ggggaagaga	aaagagaaaa	aggaggggga	aagtctgcat	aactgtcata	120
acctctgctt	ctcctgctct	gtaacaaacc	cacaaccagg	aagagtcatt	gtctggaaca	180
atcatgggac	cccaaacgcc	tgtaggtttt	ttaccacca	acatcaccca	tggtctgctt	240
aagctgtcat	ttgtttccca	cagttacctt	gcacacgga	tgcccaattt	atggcccagg	300
aaggctgacc	caggctaagg	gcagtctcac	tccacagcca	tgcaatggac	agtctgaatg	360
tttcttacct	cagaccttta	ctgacctcta	ctatttctct	ctctgatata	aaagaaaaac	420
acttttaatt	ttctnctgca	tnctacatct	cctnctaaaa	antttggcct	aattgncatc	480
aaaaccttgt	aggaatctga	aatttttggt	cttctgaatc	ttancc		526

<210> 1474

<211> 187

<212> DNA

<213> Homo sapien

<400> 1474

aaacttggtt	gctgtgaaca	attgtcgaaa	agagtcttcc	aattaatgct	ttttatatct	60
aggctacctg	ttggtttagat	tcaaggcccc	gagctgttac	cattcacaaat	aaaagcttaa	120
acacattgtc	caaaaaaaaa	aaaaaaaaaa	gccccykccc	sgggggsccck	ttmaaggggr	180
aawtccc						187

<210> 1475

<211> 474

<212> DNA

<213> Homo sapien

<400> 1475

ccattctctt	tatctcaaac	cgaagaaaga	tatgatgcag	gcagtagttt	tttcttagtg	60
cctcatagta	tctaatagca	gaaagtgcgc	gcacatagcg	agcacattag	tttttatgta	120
tctacaggac	agaagggcca	cttagctgat	ggctccaggt	ttcctttgat	ataatctaata	180
gttcttatga	cctcaaagac	tgaacacatt	tccctaagtg	cttcacttag	cacccaggag	240
caacttggag	tcttcgcaga	ataaaatcca	ttatttttaata	gtagattaata	acatgggtac	300
ttatatctat	gcaggtctat	aatagtttat	tcctatgtaa	gctttattaa	aagcattgggt	360

atgtttttaca taaaaagtta atgtgaatat tagaaaaaaa ggacaatatt aaagcagttt 420
gtagaatttg tttccccccc aaaatgaatg aaatacacaa tagatgtaca aaaa 474

<210> 1476
<211> 401
<212> DNA
<213> Homo sapien

<400> 1476
ccttggggac agggcaggag gacgcacacc tcatggacag ggcggccagg gctgagatac 60
cagcgggggtg ggtattcccc gcgggtgctt acctccaaca gtgtcttgtc agcaaaggcc 120
atgatgcctt caaagatgat gacgtttgca ccatacagtg ttttctgtga agaaaccag 180
gagttgcgga gcctggctca tgtgcctgca gcccccgag gccccctctg cagggccctg 240
gcctaccag tccttcttcc ggctgtgcgt ggtgaagtca taaatgggca ccttgacact 300
cttccccctgc ttcagcttct tgaggggtgga aatgatgaag gtcgaagtca aaaggcatct 360
gggggtgggtc gaaagtttga aagtttgctt gtgggtgccgg g 401

<210> 1477
<211> 753
<212> DNA
<213> Homo sapien

<220>
<221> misc_feature
<222> (1)...(753)
<223> n = A,T,C or G

<400> 1477
cagcatgctt aaaaagttag aggaattgga acagaaatac acctwmcaac ctkrmcctnt 60
taccaaaaac aaacnagtgg tatkggamcc sacctttmrk ctttttcmac macttatttc 120
aaagytsrnt kgtggkgaaa agmcacycyk snatscywcc rcaccttgw aggcyygttg 180
acttrataac akknctgctn atnwnrtgta ggggtgatay tgatgrtgaa attgcactta 240
gctggggttat aattkgaaag tcaaagtctt atttgataaa gatgtgaatg agagaaatac 300
agtaaaagga ttttaggaagt tcaacatttt gggcacgcac acaaaagtga tgaacatgga 360
ggagtccacc aatggcagtc tggcggctga atttcggcac ctgcaattga aagaacagaa 420
aaatgctggc accagaacga atgagggctc tctcatcggt actgaagagc ttcactccct 480
tagttttgaa acccaattgt gccagcctgg tttggtaatt gacctcgaga cgacctctct 540
gcccgttgtg gtgatctcca acgtcagcca gctcccgagc ggttgggcct ccactccttg 600
gtacaacatg ctgggtggccg gaaccagga acctgtcctt ctctctgact cccctctgtg 660
cacgatgggc tcancttttc anaagtgcct gagttggcag tttttcttnt tgtcacccaa 720
aagaaggtct caatggnngg acccanaacc ttt 753

<210> 1478
<211> 421
<212> DNA
<213> Homo sapien

<220>
<221> misc_feature
<222> (1)...(421)
<223> n = A,T,C or G

<400> 1478
aaacctatac tcactttccc aaattgaatc actgctcaca ctgctgatga tttagagtgc 60

tgtccggtgg	agatcccacc	cgaacgtctt	atctaatacat	gaaactccct	agttccttca	120
tgtaacttcc	ctgaaaaatc	taagtgtttc	ataaatttga	gagtctgtga	cccacttacc	180
ttgcatctca	caggtagaca	gtatataact	aacaaccaa	gactacatat	tgctactgac	240
acacacgtta	taatcattta	tcatatata	acatacatgc	atacactctc	aaagcaaata	300
atttttctact	tcaaaacagt	attgacttgt	ataccttgta	atttgaaata	ttttctttgt	360
taaaatagaa	tggtatcaat	aaatagacca	ttaaccaana	aaaaaaaaga	aaaaaaaaaa	420
a						421

<210> 1479

<211> 214

<212> DNA

<213> Homo sapien

<400> 1479

ggaaatatat	aataaaaaatg	ttaaccagaa	ggtaaacttg	agtgtaatg	tcagacagac	60
acacttttcc	accagtgtat	ttgaatttta	gaccagtgc	cctgttttgt	ggcattcatg	120
caaaacatgc	tgagggcttt	gttcatctgg	tcatcgtgtc	caaatttcag	tcattgtttgt	180
agcaagattt	tggaagcatt	catatttcct	tttt			214

<210> 1480

<211> 434

<212> DNA

<213> Homo sapien

<400> 1480

ggaggccgct	tacgtaaagc	ccaggggaca	ttcaacagcc	cctactacc	aggccactac	60
ccaccaaca	ttgactgcac	atggaacatt	gaggtgccc	acaaccagca	tgtgaagggtg	120
cgcttcaaat	tcttctacct	gctggagccc	ggcgtgcctg	cgggcacctg	ccccaaggac	180
tacgtggaga	tcaatgggga	gaaatactgc	ggagagaggt	cccagttcgt	cgtcaccagc	240
aacagcaaca	agatcacagt	tcgcttccac	tcagatcagt	cctacaccga	caccggcttc	300
ttagctgaat	acctctccta	cgactccagt	gacccatgcc	cggggcagtt	cacgtgccgc	360
acggggcggt	gtatccggaa	ggagctgcgc	tgtgatggct	gggccgactg	caccgaccac	420
agcgatgagc	tcaa					434

<210> 1481

<211> 131

<212> DNA

<213> Homo sapien

<400> 1481

aaaatcccca	taaatctttt	ctgtcctgag	gtagttgcaa	aataaatcat	aacttgata	60
tcaactagag	ctgaggcttt	gactttttac	tcattaaaac	tagttgttac	aggaactacc	120
tttagatatt	t					131

<210> 1482

<211> 324

<212> DNA

<213> Homo sapien

<400> 1482

tgctcgctcc	tcagaggctg	aaaacatgag	aagctagggtg	tggtgaaacc	aaagcagctt	60
tattgttcaa	atgctaaaga	cgggaggatg	gactggctca	agccttaaag	aaaccatctc	120
gactttttga	actcagtga	cgggtttaag	gaaaacgtgg	gaaatatgca	aaggtggtgc	180
aggagggtgc	aggtctgtgt	gtcttattcc	catggatatc	ttgagtaatc	gcttgtccag	240

aggtgggggtt tgtgtcatcc tgaattcaac ccagcaatgg taggggtactg ttcataactc 300
accctaagcc agaagattcc tcag 324

<210> 1483
<211> 393
<212> DNA
<213> Homo sapien

<400> 1483
atgtttaatg aatgatacag gatacatccc tgttgggaagc ttgcaaaaaga cacatacact 60
gtgggtacata tttgatttaa tagaagttgt ttatcaggct atatatatat ttgccccaac 120
atgcaccaca ggataaaata actatttaca taacataggg tatttaattg acatagacta 180
tcagctttgc tgagagcaga agatggcaaa gcaatactgc agcagaaagt ggaacaacta 240
ttctaaagca atactttaga tatatttttc tagaatggat ttattagatt actttttgga 300
aagcatttga cctaaattaa atatagagct ctgaaactta gaataaaaatt tgcacttgct 360
gaaacagaat actttgcata aaaataatcc ttt 393

<210> 1484
<211> 323
<212> DNA
<213> Homo sapien

<400> 1484
tttagatcag aaagtttgag gtcttcatca gcagacactc gtgcttctat ttttcttggt 60
ttatcgaaca gttctgaaac tttgagaaaa aacttgcata tatctgtaga atcctgagtt 120
cctaaagcat ataatgaaga accaattcta ttgtaatcat ctgcagcact tttgtgggat 180
cttgtcattc tatcagattt agcagatgca tccttaactc ggttatgata ttccaaaaga 240
aatgttcggt cgtgctcaaa gaaatcatct acatccttta ctctgaaac gattactcca 300
tctgctgatt taaccatggt ttt 323

<210> 1485
<211> 405
<212> DNA
<213> Homo sapien

<400> 1485
aggagcgtca ggaaaacacg ggcagcctgg gctctgaccc gagccactcc aactccacgg 60
ccacgcagga agaagacgag gaggaggagg agagtttttg gaccctctct gacaaatact 120
cctcccggag actattccgc aaatccgcag cccagttcca taacctgcgg tttggggaac 180
ggagagatga gcaaatggaa ccggagccca aattatggcg aggcgggaga aacaccccgt 240
actggtactt cttgcagtgc aaacacctga tcaaggaagg gaagctggtt gaagccctgg 300
acctgtttga gaggcagatg ctgaaggagg agcgattgca gcccattggag agcaactaca 360
cgggtgctgat tgggggctgc gggcggttg gctacctgaa gaagg 405

<210> 1486
<211> 230
<212> DNA
<213> Homo sapien

<400> 1486
aaaaatatgt ggattgtgct tgacgtagca aatttcttct atctgcaaaa gcccttttct 60
cactacctca tataaccccc tttgatatgg caccatgttt gaaattggag cgtacacaca 120
tagtcattgg atttactggg attctctttg tgacaagtag gagccaaggg gtcattgcagg 180
gaagcgaacg tgcccgataa ggatttcctt gttgccagag tgtttagcag 230

<210> 1487
 <211> 273
 <212> DNA
 <213> Homo sapien

<400> 1487
 tttccactct gcacattgta gagggaacac tctgtaggcc catgggtccc ttactagaga 60
 ggttgagtga atttgccttc agttaacatg ggaccttctg tttagcttcc tcttgcttcc 120
 caaagatttt aagcattttg taaatgtata aactcacctc tggtaacagt ggcccagacg 180
 ctgctttgtg ctaaaagcat gggaaatgta aaggcagtct ttctctggga aatggatgct 240
 attctattct gctgcccta cctgttcctg agg 273

<210> 1488
 <211> 452
 <212> DNA
 <213> Homo sapien

<400> 1488
 cctactgtgc cccgtaggca aagctctgaa gatttcatcg aaaaatctgc tgtcaatacg 60
 tagaaaagtt cactatttca gtttcacagc aaaaaagggtg gggggagggg ggaacccaat 120
 agatatttaa gtagatgctt tccaatccca ttcactgcat taattagctt acctcttata 180
 cagtacaaca taaacattgc atgtttatct gtatgtaaca cctataagca tatagcatct 240
 acatttttaag tgtattttaca aattcaacaa aatatctaca tataaaaagc tttactttaa 300
 attaaacttg atgcaagtta tgagaaacca atttattggc aaatgaaact gagcattcct 360
 tcaaccatag gttgttatag attttcatat ttggaggtaa cccatttgat agatattggt 420
 tatgaatacg atagaatata tatttacttt tt 452

<210> 1489
 <211> 653
 <212> DNA
 <213> Homo sapien

<220>
 <221> misc_feature
 <222> (1)...(653)
 <223> n = A,T,C or G

<400> 1489
 cctgctcttc tcttcaaagc acttagtaca cagggktaca ggtgctacca cttggattcc 60
 ccagagcatg gaagtctgat cccaggttga acatatttct tctgaaaatg agcatcttgg 120
 ttctatagat tcttatcttg ctcacaggac ttgctccaaa actgaatttt cagaagcagc 180
 atgataggga aagagatatt caactctgac agacaaggta gatcgaagca cccacactaa 240
 tttctttcag gtgccccatg aggaagactg catcatgtca ctccactca cttggggaga 300
 ttctaggact gagacacaaa gttccccag agtttctgct aatggaaggg gaaacaggtg 360
 gtttggaatg gaaaggtgga accaggtcca caaatgtgc tccctctgct caagactgac 420
 tttggctttc ccaggtcccc acttgacttt catataagct gagatgacct attacgggaa 480
 aaattaggga acacctaata aaaccaactt tcaaaaactc ctatttatca tggatgtgcc 540
 acgatcgaga gaatcnaaca cnaactgnct gtnagagagg ccttcattnt gnctcatctt 600
 gagctaaaat cctgrcttgg gatgccagaa ancatgnccc tcttntcggg ttg 653

<210> 1490
 <211> 363
 <212> DNA

<213> Homo sapien

<220>

<221> misc_feature

<222> (1)...(363)

<223> n = A,T,C or G

<400> 1490

taacctgaca	aaataaaact	tagtaaaatc	takaactggt	tcttggccta	cttgagagga	60
acttccatat	tttcacagcc	atctccgaaa	gcagcagttg	ctgtaaatta	actgagactt	120
ggaaatgggtg	cagactgtct	tggtagagct	gttcttatag	cacaatttta	tctggaaaat	180
aaacttgtaa	atgcgtgctg	tatattaata	catgtgtgoc	catatattatt	tttattatct	240
cctgccagtc	tttgctcaat	gggagatgac	agaccaactt	ctcaacgtga	tttccccatt	300
tcattgaatg	agatttatat	gccacttatg	aaaaaaaata	ctgctgngaa	agaaatgtac	360
ttt						363

<210> 1491

<211> 163

<212> DNA

<213> Homo sapien

<400> 1491

taatcagccc	ctaattttctc	catgtttaca	cttcaatctg	caggcttctt	aaagtgcacag	60
tatcccttaa	cctgccacca	gtgtccaccc	tccggccccc	gtcttgtaaa	aaggggagga	120
gaattagcca	aacactgtaa	gcttttaaga	aaaacaaagt	ttt		163

<210> 1492

<211> 184

<212> DNA

<213> Homo sapien

<400> 1492

yattccccag	gggaaaaatt	gaaagtcaaa	ctattcacca	agagaatgca	ttgtctttgc	60
aaatgagcct	aagaatcaga	ctttttataa	atacatgttc	aagtttcttg	tggttctaaa	120
tggacactga	gaactgaaac	tgtctacacc	aagtttacaa	tctatattaa	ctatcattwt	180
acag						184

<210> 1493

<211> 273

<212> DNA

<213> Homo sapien

<220>

<221> misc_feature

<222> (1)...(273)

<223> n = A,T,C or G

<400> 1493

aggtaawttg	tgatatttag	tgcacattta	cgtgtaggnc	crtcttkaat	ggtaaagaca	60
gatacaagcc	tatggcacac	ttctccaaag	caagctatac	ttgagagcca	attcccaaatt	120
aagacagcag	agatctgatt	aaatgcaact	gtgcaaacat	tcaacagaca	tgttgaatgt	180
aagacaaatt	atgattactg	ataatatgca	aatgtggtct	ataaatttat	gaatgtgact	240
tccaagggga	atatggtatg	gaagcccatt	ttt			273

<210> 1494
 <211> 343
 <212> DNA
 <213> Homo sapien

<400> 1494
 ttggaaagcc tatcactttc tctcttcatt ctccagcccc cacaccaagc acacagagct 60
 ttccagtgtc ttactcttaa tggagaacat aaccagggat tatcagggtat tccaacatga 120
 aaaagaaagt ccaatagaaa caagcaggat aatcaaacca ggaggaagca gagactatat 180
 agagaaagaa aaaaagacac atgggaataa cggcaataat actgacaata cacctacca 240
 taaacttatc agaatgaatt tgttgagaa atatatggag gggagggtact tgtgtgtgtg 300
 cacaggcact catgtacacg tgtgtatgtg tatgtttttt taa 343

<210> 1495
 <211> 378
 <212> DNA
 <213> Homo sapien

<400> 1495
 tagcattctt ccagccactc tggcgctact atgtgcttca cgacagaaat cgccgtcagg 60
 aacttcacgg tgcgagtcac tttgctggca atgaggtgtg tgcacttctg tgcagactcc 120
 gcaacctctc caccaagaat gtagagcttc ttaataact gttgaacctg gacaggctcg 180
 aatccagtga aaagcacaac aggggtcaat tctggagtta gcttttttagt gggaggtggt 240
 acgtcttcaa ttctggctct tttggaagaa ggctggacat tagctacttc attctgtttc 300
 agtttgggag gtagtcttat actcatcaac aactctgcag acacttttaa gggaactctc 360
 caagcatcta aaagattt 378

<210> 1496
 <211> 181
 <212> DNA
 <213> Homo sapien

<400> 1496
 tggagaagga agttttcctg aagagccaga atccttgcta agtcatttag atccaactga 60
 ccatctttat ttctgtcaaa aatcttcac atgggtgccag tgtattcttc cagtttagcc 120
 tcagaaatgg cctttttgtg gtgaagaaag aggtctcgga ggaagttgag gagctcagca 180
 g 181

<210> 1497
 <211> 373
 <212> DNA
 <213> Homo sapien

<400> 1497
 tggaaagctga tccaccttga gatcaagccg gccatccgga accagatcat ccgcgagctg 60
 caggctcctgc acgaatgcaa ctgcgcgtac atcggtgggt tctacggggc cttctacagt 120
 gacggggaga tcagcatttg catggaacac atggacggcg gctccctgga ccagggtgctg 180
 aaagaggcca agaggattcc cgaggagatc ctggggaaaag tcagcatcgc gggttctccgg 240
 ggcttggcgt acctccgaga gaagcaccag atcatgcacc gagatgtgaa gccctccaac 300
 atcctcgtga actctagagg ggagatcaag ctgtgtgact tcggggtgag cggccagctc 360
 atcgactcca tgg 373

<210> 1498
 <211> 337

<212> DNA

<213> Homo sapien

<400> 1498

gctctttag	tgcttttctt	ttaagggaga	tgtagtaaaa	gggaaaatgt	agctcttagt	60
ttacacttca	aagatgtggg	ggtctttcag	agaactaaga	ataacagttt	tatgtgcaga	120
gagagtttgc	cagatctgaa	gcatatacct	cattgactag	gctgttactt	tgggataggt	180
tgcagtacca	gccacagcca	gcagatagag	gaaaagacac	acataaaactc	gcttctgagc	240
gtccacttct	gcactctctg	ctctgctgtt	actcagcccc	tgagtctgac	tcattctctgc	300
acaacctctc	tgtgccatga	agataagtct	tccatgg			337

<210> 1499

<211> 314

<212> DNA

<213> Homo sapien

<400> 1499

catgctggagg	gacttttagca	tggctgataa	ggtccttctt	accattccaa	aagaacagag	60
gaccagagtt	gcacactttt	tggaaaggca	gggcttcaag	cagcaagctc	ttacagtatc	120
cacagatcct	gagcatcggt	ttgagcttgc	tcttcagctt	ggagagttaa	aaattgcata	180
ccagtttagca	gtggaagcag	agtcagaaca	gaagtggaaa	caacttgctg	aacttgccat	240
tagtaaagt	cagtttggcc	tagcccagga	gtgcctgcat	catgcacagg	attatggggg	300
cctgctgctt	ttgg					314

<210> 1500

<211> 321

<212> DNA

<213> Homo sapien

<400> 1500

cctgaaacct	ggtgggaaga	tgattgaaag	tgttttagat	tcaacagatt	gactatgtat	60
gacttatcta	ttaaaatgaa	gaacttccat	ggtttaatag	aatgaatgct	gtattcaaca	120
aggtcttcca	tccttcttat	aaatcttaag	actgtgttta	agctttcttt	cacttttact	180
ctatcccttg	gaagttaatt	gggaataaaa	agatttatca	atttagtcac	tataatttaa	240
ggccaggcat	ctgcttgga	atacaataac	cacaattaat	acttagagaa	aattgtttca	300
acagattaac	tctgctat	t				321

<210> 1501

<211> 557

<212> DNA

<213> Homo sapien

<400> 1501

ctgctctggg	gaaaatggtg	gaggagccag	gcagagagga	ggagcagagt	gctggcagtg	60
gaaagcctag	ctgagactgg	agatgcccc	ctgcccagg	catctcagcg	aggatgcttc	120
tccatattggg	tgagccagcc	tagagacaga	acaggggaag	ccagcgggtg	ctgcagcgac	180
ccaccgcccc	agaacatctg	catcttacat	caacaaaggt	ttatttctca	ttaatatcca	240
ttgtgggttg	gctgccactc	taaccctcgt	tgctctcca	tctgggtctt	gggtggcaga	300
gcagcctgtc	tctgtggcag	aggaaaagag	agcactgggc	agcacaggct	gactctcaaa	360
ttttccgcct	gaaggtgacc	caagtcactg	ctcacatttc	attgactaaa	gcaaaatcct	420
atgcctgtgg	gtgagttgag	caacgtgatg	aggtgttaac	ttcctacagg	gaggggctca	480
aatattgccc	aacagtggta	tggccactg	cctgggggtg	tcggtggaag	gctggcagga	540
caaggagagac	cacgtgg					557

```
<210> 1506
<211> 189
<212> DNA
<213> Homo sapien
```

<400> 1506
 aaaagtcata aggggttttat tttgtatcat caaaatattc tataagggtcc caaataactct 60
 ttttcaaccc atgaacagta agaattttgtg aattctgata atgaaaaaag ttttcctcca 120
 ggtatgtttg tttcacattc agtcctaaag ccttgagcta tgtgtacttc cctcacacag 180
 gaacaccag 189

<210> 1507
 <211> 268
 <212> DNA
 <213> Homo sapien

<400> 1507
 ctgcacagag gggcacggaa ctccaaatcc tggaatgagg gtcaataatg tgaattctgg 60
 ccctgaccgc cagacacaca gcaagcctga gtcactctgcc gtcaccatgt cagccacaca 120
 atcctgtccc tgggcaggct cgggtggcaat gtctgtgatt ggcatctggg gccagccag 180
 ctctctgctc agtacaatgt tgggaccctt tgctgggatg tcaaacacca gcaccgggc 240
 tgaccacgtt cccacacaga tgaagtgg 268

<210> 1508
 <211> 159
 <212> DNA
 <213> Homo sapien

<400> 1508
 aaagatggca aggcaataaa tgtgttcgta agtgccaacc gactaattca tcaaaccaac 60
 ttaatacttc agaccttcaa aactgtggcc tgaaagttgt atatgttaag agatgtactt 120
 ctcagtggca gtattgaact gcctttatct gttaaatttt 159

<210> 1509
 <211> 234
 <212> DNA
 <213> Homo sapien

<400> 1509
 ccattgtgga gtacattatg aacacaatgt gcttgykaag tcttctctct cattttcaga 60
 cagcaattgt taagagtcac acacacgtcc cagacctaag cagcaactcc agtgaatggg 120
 actcagacac actcacggga cagcacagaa cttgattctt ctttgtctgt tgcccaaaga 180
 acctgttctt tgagtctgtt ccagggtgact tgtaatgata cctcttacgg tttt 234

<210> 1510
 <211> 437
 <212> DNA
 <213> Homo sapien

<400> 1510
 aaagcagtac atcttaatat gaagacagga atttctatga tgcttacgaa cattagactc 60
 aacatttttg cagccccctt tcttgggtcta cattcacaca aacatgagac acagtcccaa 120
 gggagaaaca gatgctggag gagcatttag ggccagagtg gaggcacaga ggaagctggg 180
 atttttcaac taccctctcc ttggttactc ctgggattcc cttaggattt cacggcacia 240
 ccagcgaaga gtttgtctag attcacttcg gagtagccac ttcgggacaa gaattgctct 300
 gctgtgttct tgagttttct gtagtcctgc agaactttgg gggtaaaaaa ttgcttcttc 360
 aatttatctt tctcatgac ggtagtaagt ttctccagtg cacactccgc atcaaaaatg 420
 taccggtaaa agcacag 437

<400> 1514
ctggagatca ggaatagaac ctttccaaga tatcataata ttttctttat aggaacactg 60

agtaatggca	agaatatattt	gagctttttcc	atgggttaaga	gogatagtct	cagaggctgg	120
agaaaaatggt	cattctgctc	agtgatccag	gagtgtgagg	acagtagctt	cctttccacg	180
tccacaagac	aatgacagat	gtgttttcctt	ccttgccctt	tctagggatc	tttctagggg	240
tgttgattct	ctcacaatat	ttcaatgtcc	catttctgtg	tttcttctcc	ctccaggggc	300
tgatttacga	ttacatgagt	cttgtcacaa	taatttcctc	ctttaacatc	aaggacaagt	360
tgatcactga	gataagagct	gatagttcca	tttttattca	gtctccactt	ctgcctgaat	420
tgcccatggt	cagtccatag	agctacttta	gtcccagggtg	tgggtcccggc	cnccatcaca	480
tcaagaactg	gtttcactgg	gccttggatt	a			511

<210> 1515

<211> 176

<212> DNA

<213> Homo sapien

<400> 1515

aaaggggaag	gkgaractta	aaagtatttcc	caactagatt	atctacacca	atacattgga	60
actctatatt	ttgctttcat	tttgtcttaa	aaaaatgaaa	tagcaacgct	ctatcagtca	120
cacagaggac	atgcarattt	agcagtattg	atattatact	ctatcttgtt	ggattt	176

<210> 1516

<211> 309

<212> DNA

<213> Homo sapien

<400> 1516

ctggggaaaa	ccgtgcatta	cctgcccac	ctgttcatcg	accagctcag	caaccgcgtg	60
aaggacctga	tggtcataaa	ccgctccacc	accgagctgc	ccctcacogt	gtcctacgac	120
aagggtctac	tggggcggt	gcgcttctgg	atccacatgc	aggacacogt	gtactccctg	180
cagcagttcg	ggttttcaga	gaaagatgct	gatgagggtga	aaggaatttt	tgtagatacc	240
aacttatact	tcctggcgct	gaccttcttt	gtcgcagcgt	tccatcttct	ctttgatttc	300
ctggccttt						309

<210> 1517

<211> 182

<212> DNA

<213> Homo sapien

<400> 1517

ccaacatcta	atTTTTTTTtac	TTTTTaaatta	tagctgttgt	gactgatgtg	agatggcatc	60
ttactgtggg	TTTTgtcttg	atTTTtttat	ttgatgatta	gtaaggatga	gtgtTTTTTtc	120
atatacttga	gtgtcttctt	ttgagaaaat	atctgttcat	gtccttttgc	TTTTcttgat	180
tt						182

<210> 1518

<211> 548

<212> DNA

<213> Homo sapien

<400> 1518

cctgagggag	agggaaaagc	ggatacccac	ctgtgtcgct	gtttgcgtgc	caagtccagg	60
aacagtccat	acagccctgc	tgcacccac	gacgctgtca	caaagcagga	gttcatccga	120
ggccaagggtg	ttgtcatgag	aatattcggt	aaagtaggga	cgctgacttt	gttcttgggc	180
agattctctt	cctgtggagt	atccagcctg	tttgcttagt	tttctgttgc	ttctgggggtc	240
tgatctctat	ctgttttact	gcagtccagt	taccaaagtg	gtataagtaa	aattgaaaga	300

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<210> 1519
<211> 491
<212> DNA
<213> Homo sapien
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```
<210> 1520
<211> 169
<212> DNA
<213> Homo sapien
```

```
<210> 1521
<211> 293
<212> DNA
<213> Homo sapien
```

```
<210> 1522
<211> 386
<212> DNA
<213> Homo sapien
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1

cagatggcac	cttgaactca	tttgtaagg	gctgtctcac	tctgccagac	caacaaaaac	300
tgagactgaa	gtcgccagtc	ctgaggaagc	aggcttgccc	ccagtggaaa	cactcatttg	360
tcttcagtgg	cgtaaccca	gctcag				386

<210> 1523

<211> 178

<212> DNA

<213> Homo sapien

<400> 1523

aaaaagccta	tcccatactg	aattgtggga	acctatgaag	tgtctcttaa	tgtcaattaa	60
aagtaacagt	ggctgcagat	attgattttct	gaaagtacat	gagaatttgt	ctctaactat	120
ggttgaaaca	acaaaaccaa	atctgaatca	ggtagagggtc	taccagacac	aaactctg	178

<210> 1524

<211> 319

<212> DNA

<213> Homo sapien

<400> 1524

wycacagcwg	aaatggggca	ctgaagtgtg	gagscacaka	atgcggggagg	gcagaaccac	60
agacaggagg	ctgagattga	cctcctgagt	gcaagctggt	ctccccctca	cctcctgcac	120
cctacgcaga	tggtgcttac	cataggattg	ccgtaaaaca	gagacacgca	ccagcgagaa	180
actttagccc	ttagtateccc	atcctcagga	cagaatcact	cttaaacadg	ttgaaatata	240
tctgcttaga	gcttttctat	gtgtctatat	aatgtatgca	taatatacaa	ttagaagcat	300
gtgattttat	aacattttt					319

<210> 1525

<211> 467

<212> DNA

<213> Homo sapien

<400> 1525

ccagactaga	cagagatcag	gtcatcaggg	gagcttccga	gcttcagcaa	agcccacagg	60
tagctctgcg	aactcagaat	gctaccctac	cttccctgca	ggccgctggt	catgtctgga	120
ctcctggggg	cgctatttaa	tgtttaccct	catctccagt	gccccctcca	aggctgtgca	180
gtgtcttggg	gctctcaggg	ccaacatcga	agagatgggg	gccacctctt	aacacctggc	240
aacagtctcc	cctcatcctg	attcctgaca	acagacaaaa	caccgggtttc	tagggtttat	300
ctgtttgttt	tttgagttga	gggttcctca	gggccttggtc	attgctagtg	atgggtccct	360
ttgctgtgtg	agaacccccct	caacccccctc	ctcctccctc	tggggatgaa	gtgggagtat	420
ttggctcccc	atttttgaca	aaaggggtca	gtgcagggag	gtggagg		467

<210> 1526

<211> 439

<212> DNA

<213> Homo sapien

<400> 1526

aaactgttta	ctggagaaaa	tcctcgctca	tgtccattta	ttgttttttt	ctgtactgtg	60
atthgtttca	agcttaggaa	aactagtata	ttagagtatg	ttctaggaaa	ttaaaagatc	120
tggttagagt	aaaaagttct	ttttaagggt	cttaactaat	tttttcacaa	ctaagaaaat	180
aatgaagta	ttcttaggct	gaaattcatc	ttattttatc	ataaattaga	ttgtaggggc	240
agcctacatt	tttgtgtatg	tggtttttatt	tcttaaatga	ttgtgtgagc	ctgggtgacat	300
tttatggttc	ttgtgatcta	aactgttttt	ccaattcaca	tcttttgtcg	tgaagtgata	360

```
<210> 1527
<211> 609
<212> DNA
<213> Homo sapien
```

<400>	1527						
ctggagaact	tgggctccat	taggtgcaat	cgttggagta	attagcccat	cttttacatt		60
tcttgccaca	aaatctcgaa	gagctgccat	ttcaggttcg	gacagtgaat	acacatgtcc		120
actgggaata	ctgtgtgctc	caggtatcat	ttctatgtga	gggtcaacca	ggcgggtgatc		180
tgggtagacg	tgctcatcta	ctggagtgta	cacattctgg	acatagtaat	acctcactgg		240
ttggtaaaact	ctgtatccat	ctactggata	atagagtggc	ggttgtggtg	ctggtgggtgg		300
gagcgatggg	ggtattggag	aatacatccg	gcagtggtag	cggcagtatt	cagaatcaaa		360
gacgatagat	cgagtgtccc	atgtgatatt	gggatcatgt	gtgctcagcc	agcgaaacccc		420
taggacgaca	gggaagaatg	gagactgagt	cacatcaaat	gacagcacct	ctcgggtgatc		480
cccaggta	actatcaggt	cgtgagtttc	gtggacaact	gggcccgatg	ctatggggcg		540
cccatcaatt	gcttccacaa	gtattggacc	cgcccgggcg	gncgctcgca	agggccgaaa		600
ttccagcac							609

<400> 1528						
tgatgtaatg	aattcatatt	tattgataca	gaaaaatatg	atataatcca	tctaaaaagc	60
aagttacaaa	acagtgtaca	gtgtaccata	gtacctatga	acacaattag	tgaagtaatt	120
tgcagagcta	taataccaaa	tcagaaatta	ttttggtaat	gaatttatga	ttttcctcgt	180
tttctgattt	tttccatgat	ctcatatact	ttattctcag	aaaacaaaag	acaaaacccc	240
acacatacac	aaaaataaac	gagtaacttc	tttacaaccc	cagaggctaa	gtcagtggga	300
aaagagggaa	atgaatgggt	atgagcataa	acacagggac	aaataaaaaga	agtttgggagc	360
acagagaaca	attcacaaat	cagaaqtcac	ttt			393

```
<400> 1529
atccgataga atccagttca atgaccttca gtctttactc tgtgcaactc ttcagaatgt      60
tcttcggaaa gtgcaacatc aagatgcttt gcagatctct gatgtggtta tggcctccct     120
gttaaggatg ttccaaagca cag                                           143
```

```
<210> 1530
<211> 636
<212> DNA
<213> Homo sapien
```

```

<220>
<221> misc_feature
<222> (1)...(636)
<223> n = A,T,C or G

<400> 1530
gtggagaagc ggcttggtcg ggggtggtct cgtgggggtcc tgcctgttta gtcgctttca 60
gggttcttga gccccttcac gaccgtcacc atggaagtgt caccattgca gcctgtaaat 120
gaaaatatgc aagtcaacaa aataaagaaa aatgaagatg ctaagaaaag actgtctgtt 180
gaaagaatct atcaaaagaa aacacaattg gaacatattt tgctccgccc agacacctac 240
atttggttctg tggaattagt gaccagcaa atgtgggttt acgatgaaga tgttggcatt 300
aactataggg aagtcacttt tgttcctggn ttgtacaaaa tctttgatga gattctagtt 360
aatgctgcgg acaacaaaca aagggaccca aaaatgtctt gtattagagt ccaattgatc 420
cgaaaaacaa tttaattagt atatggaata atggaaaagg tattcctgtt gttgaacaca 480
aagctgaaaa gatgtatgtc ccmnctctca tatttgga gtccttaact tctagtaact 540
atgatgatga tgaaaagaaa gggacagggtg gtcsaatgg ctnttgagcc naattgtgta 600
acataattcag tacccaatth actgnnggaa acagcc 636

<210> 1531
<211> 194
<212> DNA
<213> Homo sapien

<400> 1531
aaaaggcaga gcattctttt ttcggcaatt ttgataagca aggtgtagat ttacattttt 60
gtccttgctc ccaacgaaat ggataaacia aaataactta ccatctactc atggaatgtt 120
gttggttag ccagtctgaa ggcccacctt aatttttata taactgtctt tagctcttct 180
tttgacaggg cagg 194

<210> 1532
<211> 300
<212> DNA
<213> Homo sapien

<400> 1532
ccatacaagg taattttgac aggttccttg gattaggaca tgggcatctt gggaggccac 60
tactggccta ccacaactgg gcagcaaaac tattacaccc tccggtataa tagttttgg 120
gtttcaatga ctggggaggaa aagggttgga attttttgct ttgggggtccc tcttaacctt 180
gtatttttaa ggtctgggac tcaccaaccc tccccttcca accagagaaa ctactgcag 240
tatctccttg aaagtctggg gacgagtctg tctaagtgct ggtgagaggc acaggaccaa 300

<210> 1533
<211> 521
<212> DNA
<213> Homo sapien

<400> 1533
gttcctttgc accctgtaga tgttctagga tagttgatgc atgttactaa attacgtatg 60
caagtctgtg agtgcgctcg aggggacatc gccaggact gactgagaca cgatgccgag 120
acctcaagcc ctgagggggc gtcccaaaac ccttacagtg aagatgttta ctcatggccc 180
ccacctctgg tccacactag aaagaagctc gcccacctc cacctgtgag atccgtgaat 240
tctcggaatg gcaggggaag ccttgcacta ggttgagag aagcatcctc cacatcctgt 300
gtcagaaaacc ctggtctccg tggcacttgt aactcacctg gctgtcttct ggtctgtgtg 360

```

```
<210> 1534
<211> 181
<212> DNA
<213> Homo sapien
```

```
<210> 1535
<211> 544
<212> DNA
<213> Homo sapien
```

```
<210> 1536
<211> 591
<212> DNA
<213> Homo sapien
```

```
<210> 1537
<211> 341
<212> DNA
<213> Homo sapien
```

```
<210> 1538
<211> 363
<212> DNA
<213> Homo sapien
```

```
<210> 1539
<211> 371
<212> DNA
<213> Homo sapien
```

```
<210> 1540
<211> 403
<212> DNA
<213> Homo sapien
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```
<210> 1541
<211> 428
<212> DNA
<213> Homo sapien
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<400> 1541
 taaaacaaaa ctaaagaaga gaaaatatat tctcgtaaat tatctgaact taaaagatgg 60
 aagcctggag atagatttgt gataagccat tgctgagtac atcctagagt tcttgataat 120
 ttcagttggg taaattacaa tagtttgcta tttcctccct cacattttat gttctacagt 180
 atctagctgc ttgggttttc ctgtatacca tggggcttct gtcattctggg ctttactcag 240
 tggcatattc cctctgccta aaactctcct cccctctcca ccttagaagt agcttttctc 300
 agaacgggtt tcccagggtt tcacctaagg tgatagtaca atctacaggg acctgcacat 360
 gaagaccttt gcatacatgc caggaagttg gactttatct ttggaaaaag ggagcctttg 420
 aagggtttt 428

<210> 1542
 <211> 345
 <212> DNA
 <213> Homo sapien

<400> 1542
 awttaaatgc ttagcaagca gcaattccac gatgggtcaaa ttcctaatat gagagaagta 60
 gaaataggaa aaataggtca ccctgatact tatgttttca ttttgcttaa tatacgtttg 120
 tatatttcaa tataacatta atagatatcg tgtcccttca cagttctaaa gtagtaagca 180
 aaatgaatta atttaaccta tgcaattaaa accaatttgg aagaatattg aggtagcaca 240
 ctgttacggg aattagtagt actcagtaat gcagttgaaa gttagtggct cctaattccag 300
 tatgaatcat ggagatgaga gaaatgatta gataaagaga tatatt 345

<210> 1543
 <211> 420
 <212> DNA
 <213> Homo sapien

<400> 1543
 aatattgaat ttctagaagc agtatattgc ttactgcttc ttaattacgt tatagatgag 60
 gtggaaatga taaaaactaa agaagcaaga ttaatcttta acacacattt caggctgttg 120
 taaaagaata aacaatgctt catataaact tctagcaaat gacttcctaa tgagggtcttg 180
 aaacagtctt tagggcacgg aatgtcatca cataattaag cagctttaag cctttattaa 240
 aaggcttaaa gtcgcaaaca atgaaatctg aaacaaactg taccatatta aactttttga 300
 tgatattttca aattcagtaa aagaaaaaaa ggatgggttca gaataacatc acgtattcta 360
 atcctgaaac acataacaaa tgcattctgaa acagcaattc ttaaaaaggt tttgcccttt 420

<210> 1544
 <211> 306
 <212> DNA
 <213> Homo sapien

<400> 1544
 ctggcttcac tctactccc tctctgctcg cagcaagtcg gccgccagct ctttgatgtg 60
 ttcccaggcc cgctgcacat gggcagattc caccgtgcga gaacagatgg caaagcgcag 120
 gacaaacttg tcctgaggt gacatggaac caagtggatt tttttggcac tgtttattct 180
 ttgcagaaga gcttcattca ctttgttggg acccttttagc cgaaagcaga caagccccag 240
 aatgacttcc acacagattt caaagcgggg atcctggcgc accagtgact caaactcatg 300
 ggacag 306

<210> 1545
 <211> 110
 <212> DNA

<213> Homo sapien

<400> 1545

ctgtctcggg	ccttcatacct	gaagatcagc	gtgtgcgatg	ccgtcctgga	ccacaacccc	60
ccaggctgta	ccttcacagt	cctgggtgcac	acgagagaag	ccgccactcg		110

<210> 1546

<211> 239

<212> DNA

<213> Homo sapien

<400> 1546

aaagaaatat	gacacgggtgt	tggatattct	aagagacttt	tttgaactca	gacttaaata	60
ttatggatta	agaaaagaat	ggctcctagg	aatgcttggt	gctgaatctg	ctaaactgaa	120
taatcaggct	cgctttatct	tagagaaaat	agatggcaaa	ataatcattg	aaaataagcc	180
taagaaagaa	ttaattaaag	ttctgattca	gaggggatat	gattcggatc	ctgtgaagg	239

<210> 1547

<211> 527

<212> DNA

<213> Homo sapien

<400> 1547

aaaaattcca	gttgagattt	ttctggttct	ctgtataaag	attgactgga	acatatacat	60
tttgggggtt	atgtttggag	actttggctc	ttattcaaac	cttccatttt	agttggcttc	120
ttctgacagt	gcttcagcat	ggaagcaagg	agggggcctc	attactgcca	ggtaagggtta	180
aaaatctagt	ttctctgctg	ggtctccatt	gtcactaaga	aaggaatggc	tctgttattg	240
ctgggcaggg	ttggctgttc	caactgataa	tcctatgtct	gggagggcta	ggagtgcctc	300
cttgctgttc	ctcttgttgt	ttccactgac	agtggagtgg	ccttgttact	gctgggtggg	360
ggttgagagt	tctggctctc	tactaggagg	gacacaacct	cagtgtagag	aggcggggat	420
accttgttac	tgtcaggcac	aggcggagg	ccagtctcct	tactccacct	acccaacagg	480
gtagcttgag	gcacttcatt	attgcctagt	gagagtggaa	gttttagg		527

<210> 1548

<211> 333

<212> DNA

<213> Homo sapien

<400> 1548

ctgtgggcgg	agctagtagg	ggcggggcta	cgtgattgac	acttctctcc	tcagacttca	60
agggtacca	ctggaccctt	cccctgtctt	gaaccctgag	ccggcaccat	gcacggacgc	120
ctgaagggtg	agacgtcaga	agagcaggcg	gaggccaaaa	ggctagagcg	agagcagaag	180
ctgaagctat	accagtcagc	caccagggcc	gtattccaga	agcgccaggc	tggtagctg	240
gatgagtcg	tgctggaact	gacaagccag	attctgggag	ccaaccctga	ttttgccacc	300
ctctggaact	gccgacgaga	ggtgctccag	cag			333

<210> 1549

<211> 438

<212> DNA

<213> Homo sapien

<400> 1549

ttgacagtgt	acgctggagc	aggttccagg	gtggggctgc	cctgccgcct	gcctgctggg	60
gtggggaccc	ggtctttcct	cactgccaag	tggactcctc	ctgggggagg	ccctgacctc	120

ctggtgactg	gagacaatgg	cgactttacc	cttcgactag	aggatgtgag	ccaggcccag	180
gctgggacct	acacctgcca	tatccatctg	caggaacagc	agctcaatgc	cactgtcaca	240
ttggcaatca	tcacagtgc	tcccaaattc	tttgggtcac	ctggatccct	ggggaagctg	300
ctttgtgagg	tgactccagt	atctggacaa	gaacgctttg	tgtggagctc	tctggacacc	360
ccatcccaga	ggagtttctc	aggaccttgg	ctggaggcac	aggaggccca	gctcctttcc	420
cagccttggc	aatgccag					438

<210> 1550

<211> 204

<212> DNA

<213> Homo sapien

<400> 1550

aaaactaagt	tattccaaca	ctaaaagcat	acaacagcat	gccaacagta	atatattatt	60
ctccaagact	ttacctatgt	aagtgttcaa	aactctgcag	cattaaacaa	cgtgtatgca	120
aattgttatg	gatacatctt	agaatctaag	aaatcaggca	agtgtttaa	aggccaacgg	180
tccaagggat	tacatctgca	gttt				204

<210> 1551

<211> 132

<212> DNA

<213> Homo sapien

<400> 1551

ccatctgtgg	atttgtctgt	gcacctattg	gctcttctag	ctgactcttc	tggttgggct	60
tagagtctgc	ctgtttctgc	tagctccgtg	tttagtccac	ttgggtcacc	agctctgcca	120
agctgagcct	gg					132

<210> 1552

<211> 433

<212> DNA

<213> Homo sapien

<400> 1552

ctgaatagag	gtcaacacag	ttgcgatgtt	gagggatggt	ctccaagcac	cttttgggtg	60
caatttgaga	acatccagac	aaatccttcc	agcagaatca	atgtttggat	gataaattgg	120
agtgagaaat	cggatctgag	gaggttcaaa	tgggtacctc	tcaggaatga	taacttctag	180
cttaaaaaaca	cctttctcat	aagggtgtgt	ggctccacct	aatatttgag	ctcgcaggct	240
atccatttgg	tctttatctt	gccaacatgt	gatgcctggg	ggtggctctg	tggctaaccat	300
gtgcagctct	ctcttcagac	gtgaagctct	ctgcatgatc	cccaagtaga	aggaaccaca	360
cacagttcac	tgctccacac	taagagctgs	ctgggatgca	ctgagctgac	acccttcaca	420
acgcagcaac	gcg					433

<210> 1553

<211> 316

<212> DNA

<213> Homo sapien

<400> 1553

gagcaaggtc	tgctgagaac	agacccagtc	cctgaggaag	gagaagatgt	tgctgccacg	60
atcagtgcc	cagagaccct	ctcggaagag	gagcaggaag	agctaagaag	agaacttgca	120
aaggtagaag	aagaaatcca	gactctgtct	caagtgttag	cagcaaaaaga	gaagcatcta	180
gcagagatca	agcggaact	tggaaatcaat	tctctacagg	aactaaaaca	gaacattgcc	240
aaaggggtggc	aagacgtgac	agcaacatct	gcttacaaga	agacatctga	aaccttatcc	300

caggctggac agaagg

316

<210> 1554

<211> 542

<212> DNA

<213> Homo sapien

<220>

<221> misc_feature

<222> (1)...(542)

<223> n = A,T,C or G

<400> 1554

aaaggaatta	ttctggcagc	acatgtagta	ttcttggaatg	atcttgctgc	tcttatttct	60
ccttttgtgt	gtgtgtgtgt	gtgtgtggct	atgggttttc	atttgtaact	ccatctgctt	120
argagagtgg	gctctctata	agggaacctg	ctgtaaaactt	cattgcagca	aggatgtaga	180
gagaaatagg	acttaattcc	actaggggct	ctcatctcac	accttaagga	ggagatttct	240
agaaaaactg	ggccagatth	tctttgytct	ccatcatttt	aatgtggcag	gctgytcagt	300
tttcttactc	ttacctatgw	gatatttctt	cgtaacgtgt	ccaaaaagaa	aaaagaccca	360
atcagtgtct	cttgactttg	ttctttgatc	cctcagtttc	ttcttgattt	cagcatgtgt	420
ccgggttcct	aattttgggt	atgagttagc	aaatttaacc	attgtgtttg	tgccctaccc	480
aggggactcc	ccagtttctg	acttgaagta	gactganaag	aatccacgag	gngctatttt	540
gg						542

<210> 1555

<211> 117

<212> DNA

<213> Homo sapien

<400> 1555

ctgtctgtgg	cttcccatgt	ctttctccaa	agttatccag	agggttggtga	ttttgtctgc	60
ttagtatctc	atcaacaaag	aaatattatt	tgctaattaa	aaagttaatc	ttcatgg	117

<210> 1556

<211> 111

<212> DNA

<213> Homo sapien

<400> 1556

ctgctgcagc	cgcagtttct	catccggagt	gtaccccgctc	atgtcgccgc	tggtaccaac	60
gcaaaaggac	acggcgccacc	ctcgaactac	ggactagtta	cttaagcgcg	c	111

<210> 1557

<211> 454

<212> DNA

<213> Homo sapien

<400> 1557

cgaggactga	tcctctagta	ctaagtgact	ggggatatta	caytarccaa	cattgggttga	60
tacatacctk	artmatcatw	tgaggaygca	gtgataarsg	satawwmywg	tatsatccya	120
acaygyacta	rctcaaaaac	tagtgggggc	ggattgatct	cctgtggggac	wkcacatgsc	180
ctgaaagtga	acatgmtcmt	ratcacctgc	agrgcttgag	atggyccmca	tkgcwgcact	240
ccgccccyac	akttttttgaw	tcwacwggag	ttaggswgmt	yctwgawtta	kcctttctac	300
ctgcctccyg	akagrwcwc	wygastwga	kgaatssatt	gackkctaag	rttakacttc	360

```
<210> 1558
<211> 404
<212> DNA
<213> Homo sapien
```

```
<210> 1559
<211> 266
<212> DNA
<213> Homo sapien
```

```
<210> 1560
<211> 142
<212> DNA
<213> Homo sapien
```

```
<210> 1561
<211> 381
<212> DNA
<213> Homo sapien
```

<210> 1562

<211> 368
 <212> DNA
 <213> Homo sapien

<400> 1562
 ggagaaagga gaaccgtaca tgagcattca gcctgctgaa gatccagatg attatgatga 60
 tggcttttca atgaagcata cagccaccgc cggtttccag agaaaccacc gcctcatcag 120
 tgaaattctt agtgagagtg tggtgccaga cgttcgggtca gttgtcacia cagctagaat 180
 gcaggtcctc aaacggcagg tccagtcctt aatgggtcat cagcgaaaac tagaagctga 240
 acttcttcaa atagaggaac gacaccagga gaagaagagg aaattcctgg aaagcacaga 300
 ttcatttaac aatgaactta aaagggttggtg cggctctgaaa gtagaagtgg atatggagaa 360
 aattgcag 368

<210> 1563
 <211> 411
 <212> DNA
 <213> Homo sapien

<220>
 <221> misc_feature
 <222> (1)...(411)
 <223> n = A,T,C or G

<400> 1563
 accwtrsaac tgcawttatt acctatgcta gntttggata agaamtgkyc wtayatgtga 60
 kagcaagagg gcacyaraws wrcttsaaca ccaawgggcm ktactwtata kawmcgawgg 120
 gcatgctwtm atgaccaact grmtgactgt ttgagaatgg acaargtgct agcgctaaac 180
 ctgtccttct tgaacrtggc ttgactaacg kcwttgatac gtttccttca kkasaataact 240
 attactasac tttgktgctt gattaccgac tgggtgactc ttgmtctcac ctatgargac 300
 agtgcctttac acaaactcrt akggaaaatt gnntttgtmc tgtganctac tcatcygaga 360
 nctccctaag ggctaacatt ncatgtttcc gtctcactag ctacacgttc t 411

<210> 1564
 <211> 602
 <212> DNA
 <213> Homo sapiens

<220>
 <221> misc_feature
 <222> (1)...(602)
 <223> n = A,T,C or G

<400> 1564
 ctagtttttaa gatcagagtt cacttttcttt ggactctgcc tatattttct tacctgaact 60
 tttgcaagtt ttcaggtaaa cctcagctca ggactgctat ttagctcctc ttaagaagat 120
 taaaagagaa aaaaaaaggc ctttttaaaa atagtataca cttatttttaa gtgaaaagca 180
 gagaattttta tttatagcta attttagcta tctgtaacca agatggatgc aaagaggcta 240
 gtgcctcaga gagaactgta cgggggtttgt gactggaaaa agttacgttc ccatttctaat 300
 taatgccctt tcttatttta aaacaaaacc aaatgatatc taagtagttc tcagcaataa 360
 taataatgac gataatactt cttttccaca tctcattgtc actgacattt aatgggtactg 420
 tatattactt aatttattga agattattat ttatgtctta ttaggacact atgggttataa 480
 actgtgttta agcctacaat cattgatttt tttttgttat gtcacaatca gtatattttc 540

tttgggggta cctctctgaa tattatgtaa acaatccaaa gaaatgattg tattaannat 600
tt 602

<210> 1565
<211> 473
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (1)...(473)
<223> n = A,T,C or G

<400> 1565
ctagtccagt gtgggtggaat tcatccaggg ggctacccct ggctctctgt tgccagtggg 60
catcatcgca gtgggtgtct tcctcttcct ggtggctttt gtgggtgtgt gcggggcctg 120
caaggagaac tattgtctta tgatcacgtt tgccatcttt ctgtctctta tcatgttggt 180
ggaggtggcc gcagccattg ctggctatgt gtnagagat aaggatgatg cagagtttaa 240
taacaacttc cggcagcaga tggagaatta cccgaaaaac aaccacactg nttnatcct 300
ggacaggatg caggcagatt ttaagtgtct tggggctgct aactncacag attgggagaa 360
aatcccttcc atgtngaaga accgagtcct cgactcctgc tgcattaatg ttactgtggg 420
ctgtgggatt aatttcaacg anaaggcgat ccataaggag ggctgtgtgg aga 473

<210> 1566
<211> 53
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (1)...(53)
<223> n = A,T,C or G

<400> 1566
ctagttatta atagnaatca attncggngt cattagttca tagcccatat atg 53

<210> 1567
<211> 136
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> (1)...(136)
<223> n = A,T,C or G

<400> 1567
ttattgattt ttttttttca ctttcccat cacaactcaca cgcacgctca caactttttat 60
ttgccataat gaaccgtcca gccctgtgg ngatctccta tganaacatg cgtttttntga 120
taactnaca ccctac 136

<210> 1568
<211> 192
<212> DNA

<220>

 $\langle 222 \rangle \quad (1) \dots (192)$

<223> n = A, T, C or G

<400> 1568

tttgngtctgt	gtgagngngt	tgaccttccct	ccatccccctg	gtccttcnct	tncccttnccg	60
aggcacagag	agacagggca	gnatccacgt	ncccatntng	gaggcagana	aaagagaaaag	120
tgntttatat	acgggtactta	tttaatatcc	ntttntaatt	anaaaantnaa	acagttaatt	180
taattaaaqa	qt					192

<210> 1569

<211> 575

<212> DNA

<213> Homo sapiens

<220>

<221> misc feature

 $\langle 222 \rangle \quad (1) \dots (575)$

<223> n = A, T, C or G

<400> 1569

ctagtgtctgt	ccccccagga	gacctgggttg	tgtctgtgtg	agtggttgac	cttcctccat	60
cccctgggtcc	ttcccttccc	ttcccagggc	acagagagac	agggcaggat	ccacgtgcc	120
attgtggagg	cagagaaaag	agaaagtgtt	ttatatacgg	tacttattta	atatcccttt	180
ttaattagaa	attaaaacag	ttaattttaat	taaagagtag	ggtttttttt	cagtattctt	240
ggttaatat	taatttcaac	tatttatgag	atgtatcttt	tgctctctct	tgctctctta	300
tttgtaccgg	tttttgtata	taaaattcat	gtttccaatc	tctctctccc	tgatcggnga	360
cagtcactag	cttatcttga	acagatatatt	aattttgcta	acactcagct	ctgccctccc	420
cgatcccctg	gtcctccagc	acacattcct	ttgaaataag	gtttcaatat	acatctacat	480
actatatata	tatttggcaa	cttgnatttg	ngngtatata	tatatatata	tgtttatgta	540
tatatgnqat	tctgataaaa	taqacattgc	tatttc			575

<210> 1570

<211> 392

<212> DNA

<213> Homo sapiens

<220>

<221> misc feature

 $\langle 222 \rangle \quad (1) \dots (392)$

<223> n = A, T, C or G

<400> 1570

ctagtcacagn	gtggtggaat	tccgcgcgcca	tcatgggtcg	catgcatgct	cccggaagg	60
gcctgtccca	gtcggcttta	ccctatcgac	gcagcgtccc	cacttggttg	aagntgacat	120
ctgacgacgt	gaaggagcag	atttacaac	tggccaagaa	gggccttact	ccttcacaga	180
tcggtgtaat	cctgagagat	tcacatgggtg	ttgcacaagt	acgttttgtg	acaggcaata	240
aaattttaag	aattcttaag	tctaagggaac	ttgctcctga	tcttcctgaa	gatctctacc	300
atttaattaa	gaaagcagtt	gctgttcgaa	agcatcttga	gaggaacaga	aaggataagg	360
atgctaatt	ccqncqtatt	cttaataqaa	gc			392

<210> 1571
 <211> 390
 <212> DNA
 <213> Homo sapiens

```
<400> 1571
gaaggacgtt tgtgttgga gccctggtat ccccggaact cctggatccc acggcctgcc 60
aggcaggac gggagagatg gtgtcaaagg agaccctggc cctccggggc ccatgggtcc 120
acctggagaa atgcatgtc ctctggaaa tgatgggctg cctggagccc ctggtatccc 180
tgagagtggt ggagagaagg gggagcctgg cgagaggggc cctccagggc ttccagctca 240
tctagatgag gagtccaag ccacactcca cgactttaga catcaaatac tgcagacaag 300
gggagccctc agtctgcagg gctccataat gacagtagga gagaaggctt tctccagcaa 360
tgggcagtc atcacttttg atgccattca                               390
```

<210> 1572
 <211> 383
 <212> DNA
 <213> Homo sapiens

<220>
 <221> misc_feature
 <222> (1)...(383)
 <223> n = A,T,C or G

```
<400> 1572
ctgcagcttc tgctgctgag gccgggattg ctacgactgg gactgaaggt gaaagaggtg 60
gaaatccgaag tcctgggact gcgggatgct aaacattgaa agctgggtgt aggcactgca 120
gggagagtgt ggaggtctga cagggtagga atatgtggga gggctgggct aggaatggcc 180
ttggaggctg gcctgtgtgg atatggcacc aattctaccc tgctcctctt ttcttttccc 240
cagactcaga cgatgccttg ctgaagatga ccatcagcca gcaagagttt ggccgactg 300
ggcttctcta cctaagcagt atgactgagg aagagcagat tgcttatgcc atgcagatgt 360
ccctgcangg gagcagagtt tgg                               383
```

<210> 1573
 <211> 149
 <212> DNA
 <213> Homo sapiens

```
<400> 1573
cctccagagc ctctctagtg gcagagcagc tcacactccc tccgctggga acgatggctt 60
ctgcctagta cctatccttg tgtttctgat gcagtggtag cattgggttca agttctctcc 120
tgctgtggtc agagttgctt cgatgttg                               149
```

<210> 1574
 <211> 143
 <212> DNA
 <213> Homo sapiens

```
<400> 1574
ctgccaggct gaaaagaagc ctgagctccc acaccgcctt cctcacggcc ctctctcggg 60
agtcaattcc actggtggac cacgggcccc cagccctgtg tgggccttgt ctgtctcagc 120
tcaaccacag tctgacacca gag                               143
```

<210> 1575

<211> 112
 <212> DNA
 <213> Homo sapiens

<400> 1575
 ctgcatccac cctctttcag ggggtagagc cactatactt ctcatgtaga tcagccacat 60
 tgtcactgga gactcggatc cagccatcct cccgcacgtg gtagagggtg ac 112

<210> 1576
 <211> 198
 <212> DNA
 <213> Homo sapiens

<400> 1576
 ccagtatgtc cccaggatta tgtttgttga cccatctctg acagtttagag ccgatatcac 60
 tggaagatat tcaaatcgtc tctatgctta cgaacctgca gatacagctc tgttgcttga 120
 caacatgaag aaagctctca agttgctgaa gactgaattg taaagaaaaa aaatctccag 180
 gcccttctgt ctgtcagg 198

<210> 1577
 <211> 444
 <212> DNA
 <213> Homo sapiens

<400> 1577
 cctgcctgga gccccagatc accccttctt actacaccac ttctgacgct gtcattttcca 60
 ctgagaccgt cttcattgtg gagatctccc tgacatgcaa gaacagggtc cagaacatgg 120
 ctctctatgc tgacgtcggg ggaaaacaat tccctgtcac tcgaggccag gatgtggggc 180
 gtcacacagg gtccctggagc ctggaccaca agagcgccca cgcaggcacc tatgaggtta 240
 gattcttctga cgaggagtcc tacagcctcc tcaggaaggc tcagaggaat aacgaggaca 300
 tttccatcat cccgcctctg ttacagtca gcgtggacca tcggggcact tggaaacgggc 360
 cctgggtgtc cactgagggtg ctggctgcgg cgatcggcct tgtgatctac tacttggcct 420
 tcagtgcgaa gagccacatc cagg 444

<210> 1578
 <211> 294
 <212> DNA
 <213> Homo sapiens

<400> 1578
 ccacaaagcc attgtatgta gcttttagctc agcgcaaaga agagcgccag gctcacctca 60
 ctaaccagta tatgcagaga atggcaagtg tacgagctgt gccaaccct gtaatcaacc 120
 cctaccagcc agcacctcct tcagggtact tcatggcagc tatcccacag actcagaacc 180
 gtgctgcata ctatcttcct agccaaattg ctcaactaag accaagtccc cgctggactg 240
 ctcagggtgc cagacctcat ccattccaaa atatgccggg tgctatccgc ccag 294

<210> 1579
 <211> 295
 <212> DNA
 <213> Homo sapiens

<220>
 <221> misc_feature
 <222> (1)...(295)

<223> n = A,T,C or G

<400> 1579

```
ccacaaaagcc attgtatgta gcttttagctc agcgcaaaga agagcgccag gctcacctca 60
ctaaccagta tatgcagaga atggcaagtg tacgagctgt gcccaaccct gtaatcaacc 120
cctaccagcc agcacctcct tcaggttact tcatggcagc tatcccacag actcanaacc 180
nngctgcata ctatcctcct agccaaattg ctcaactaag accaagtccc cgctggactg 240
ctcagggngc cagacctcat ccattccaaa aatatgcccc gtgctatccg cccag      295
```

<210> 1580

<211> 166

<212> DNA

<213> Homo sapiens

<400> 1580

```
cttcttttatt ggggacatgt gggctggaac agcagatttc agctacatat atgaacaaat 60
cctttattat tattataatt atttttttgc gtgaaagtgt tacatattct ttcacttgta 120
tgtacagaga gggtttttctg aatatattatt ttaagggtta aatcac      166
```

<210> 1581

<211> 449

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (1)...(449)

<223> n = A,T,C or G

<400> 1581

```
ctgaggcaac agaataaatg cagaggcatt acaatgaatc ccaacttaata taaagaacta 60
tacagaccaa cacttctcta caaaatTTTT ttttcctcat tgccagttaa atacagagtt 120
ttactttcat agcttaacaa tgaagggtca taaactgaag ccaatacata tacctagcat 180
ttcagtctaa gcttgtccac gtacatagct gaagtcaatt acaaggtttg gcctagaaat 240
gctaggggaa cttcttttga gttttttacag gtattaaact tcatcttgca cactgaagtc 300
atcatacata cagggcaaaa tcagagcttt tatatttgcg tttattcttc atttaacttt 360
ttataacact actatagttt attaaaacaa aaaacaaaga gcaagtagtg agcatattan 420
gattacagtc ctttctactca ttcacacct      449
```

<210> 1582

<211> 302

<212> DNA

<213> Homo sapiens

<400> 1582

```
ccaatgggct ttgctgtagc ttgctgaaat caccaagcag gagagattta accagaggcg 60
atgtgtccag tcaccagcat agagccatcc tctgtgtcac catccacacg cagggccttc 120
tggcagacct catgcaatgc cctccatgtt aatattcatc agaaaatgga taattagggg 180
ggccagcaaa aatatcaagg gtcaaataac gcacatttct gtttaggcca tctatggctt 240
tcatctcttc tgaagtcaac tggaattcaa acacctgcac gttctgtctg atgcgctgct 300
ca      302
```

<210> 1583

<211> 170

```
ccaatgtaca tgggtggacta tgccggcctg aacgtgcagc tcccgggacc tcttaattac 60
taaacctcag tactgaatca ggacc                                     85
```

<210> 1588
 <211> 369
 <212> DNA
 <213> Homo sapiens

<220>
 <221> misc_feature
 <222> (1)...(369)
 <223> n = A,T,C or G

```
<400> 1588
ccaggctacc ttcccactgg agacaggcag ggggacaggt gctaagggac ctggcaggca 60
gggctggcag gccccatggc gcctgttcca gcagatgaca agcccaggtc agggtagagc 120
gggcaggagg ggggacgagg gctcccacaa catgattttg tgtaaaatat ggcagcgaca 180
cacgctcagg gccgggaggt ggggggttagg gtggggacgg cggcaacatc gtgtaaaaaa 240
gtgtcccagt tcccatagca aagagagctg tgaccgggtg ttcagagctt ctccagtaca 300
agggggaaag ccgcccggcg ggggcggcgg gcaggggacat catttggttt cctggtgctg 360
tcngtccga                                     369
```

<210> 1589
 <211> 361
 <212> DNA
 <213> Homo sapiens

```
<400> 1589
ctgtagcttc tgtgggactt ccactgctca ggcgtcaggc tcagatagct gctggccgcg 60
tacttgttgt tgctttgttt ggaggggtgtg gtggtctcca ctccgcctt gacggggctg 120
ctatctgcct tccaggccac tgtcacggct cccgggtaga agtcacttat gagacacacc 180
agtgtggcct tgttggcttg aagctcctca gaggagggcg ggaacagagt gaccgagggg 240
gcagccttgg gctgaccagc gacggtcagc ttggtccctc cgccgaacag taaaaagga 300
ctcaggctgt tatcatagga ctggcagtaa taatcagcct catcttcagc ctggagccca 360
g                                             361
```

<210> 1590
 <211> 434
 <212> DNA
 <213> Homo sapiens

<220>
 <221> misc_feature
 <222> (1)...(434)
 <223> n = A,T,C or G

```
<400> 1590
ctggagaagg tgtgcagggg aaaccctgct gatgtcacgg aggccagggt gtotttctac 60
tcgggacact ctcccttttg gatgtactgc atggtgttct tgggtgctgta tgtgcaggca 120
cgactctgtt ggaagtgggc acggctgctg cgaccacag tccagttctt cctggtggcc 180
tttgccctct acgtgggcta caccgcgtg tctgattaca aacaccactg gagcgatgtc 240
cttgttggcc tcctgcaggg ggcactggtg gctgccctca ctgtctgcta catctcagac 300
ttcttcaaag cccgaccccc acagcactgt ctgaaggagg aggagctgga acggaagccc 360
agcctgtcac tgacgttgac cctgggcgag gctgacnaca accactatgg ataccgcac 420
tcctcctcct gagg                                     434
```

<210> 1591
 <211> 439
 <212> DNA
 <213> Homo sapiens

<220>
 <221> misc_feature
 <222> (1)...(439)
 <223> n = A,T,C or G

<400> 1591
 gctttcgcca gaaaatgttg catgtcaaac aatatgtgat ccatactgtg tgcgctcctt 60
 gggggtttat ttgactttgt cacaatgaca gccaacagtg agactgataa gcctgtaaaa 120
 ataaaaaaat aagactaatc aaatagacat ggcattttta tctcaaagtg caaaatcatc 180
 taactgaaaa tgacggcatt gagaaattcc agtgggttaa aatgaatcaa aacttcatta 240
 cgcaggcagt ggaagtgtgt tgaaagattt accaggggtg tcaagtttta gacactcaga 300
 aaggcaccat tctagccatc ttgattggat aacatgtata tacttatgtc cctacgatat 360
 tcaaaagata atactgtttt agtacaaaac aatcaaaca ggcaaagant caaaaccaag 420
 ccaaccctaa tatccccag 439

<210> 1592
 <211> 74
 <212> DNA
 <213> Homo sapiens

<220>
 <221> misc_feature
 <222> (1)...(74)
 <223> n = A,T,C or G

<400> 1592
 tttttttttt taatgttcac agtccctgct ttatttccat ttgttcacac acncttttaa 60
 aaaaaaaaaa aaaa 74

<210> 1593
 <211> 288
 <212> DNA
 <213> Homo sapiens

<400> 1593
 ccattccgaag caagattgca gatggcagtg tgaagagaga agacatattc tacacttcaa 60
 agcttttggtg caattcccat cgaccagagt tgggtccgacc agccttggaa aggtcactga 120
 aaaatcttca attggattat gttgacctct accttattca tttccagtg tctgtaaagc 180
 caggtgagga agtgatccca aaagatgaaa atggaaaaat actatttgac acagtggatc 240
 tctgtgccac gtgggaggcc gtggagaagt gtaaagatgc aggattgg 288

<210> 1594
 <211> 455
 <212> DNA
 <213> Homo sapiens

<400> 1594
 ccacacagac tcaccaagcc acagacttgt cttccacaag cacgttctta ccttagccac 60
 gaagtgacca agccacagct actaaagggt gaactcaaa atagtgtacag ggtattaaac 120

<210> 1595

<212> DNA

<220>

 $\langle 222 \rangle \quad (1) \dots (367)$

<400> 1595

<210> 1596

<212> DNA

<400> 1596

<210> 1597

<212> DNA

<400> 1597

<210> 1598

<212> DNA

<400> 1598

ctgcctataa aactagactt ctgacgctgg gctccagctt cattctcaca ggatcatcatc 60

```
ctcatccggg agagcagttg tctgagcaac ctctaagtcg tgctcatact gtgctgccaa 120
agctgggtcc atgacaactt ctggtggggc gagagcaggc atggcaacaa atcccaagtt 180
agggctcca atgagcttcc tagcaagcca gaggaagggc ttttcaaagt tgtagttact 240
tttggcagaa atgtcgtagt actgaagatt cttctttcgg tggaagacaa tggatttcgc 300
cttcactttc ctgtccttaa tatccacttt gttgccacac aacacaatgg ggatgttttc 360
acacactcgt accagatctc tatgccagtt aggcacattc ttgtaagtaa ctctcgatgt 420
tacatcaaac attatgatgg cacac 445
```

<210> 1599

<211> 142

<212> DNA

<213> Homo sapiens

<400> 1599

```
cctgccccag ggggaagcac ggacccgaga cgacggcgat gaggaagggc tcctgacaca 60
cagcgaggaa gagctggaac acagccagga cacagacgag gatgatgggg ccttgacagta 120
agcagcctga caggagcaat gg 142
```

<210> 1600

<211> 297

<212> DNA

<213> Homo sapiens

<400> 1600

```
cctgcacttg aacatggctt tggttttaag caacttctct accctgaccc tcctcctggg 60
acagcgtttc gggaggtttc ttggcctcac tgagagggat gtggagctgc tgtaccccgt 120
caaggagaag gtattctaca gcctgatgag ggagagcggc tacatgcaca tccagtgcac 180
caagcctgac accgtaggct ctgctctgaa tgactctcct gtgggtctgg ctgcctatat 240
tctagagaag ttttccacct ggaccaatac ggaattccga tacctggagg atggagg 297
```

<210> 1601

<211> 289

<212> DNA

<213> Homo sapiens

<400> 1601

```
ctggagatga tcctcaacaa gccagggctc aagtacaagc ctgtctgcaa ccaggtggaa 60
tgtcatcctt acttcaacca gagaaaactg ctggatttct gcaagtcaaa agacattgtt 120
ctggttgctt atagtgtctt gggatcccac cgagaagaac catgggtgga cccgaactcc 180
ccggtgctct tggaggaccc agtcctttgt gcctcggcaa aaaagcacia gcgaacccca 240
gccctgattg ccctgcgcta ccagctacag cgtgggggtg tggctcctgg 289
```

<210> 1602

<211> 398

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (1)...(398)

<223> n = A,T,C or G

<400> 1602

```
gggagggcag agggagaatg ggaagatcag gaagctctag attacttcag tgataaagag 60
```

<210> 1603

<211> 438

<212> DNA

<213> Homo sapiens

<400> 1603

<210> 1604

<211> 297

<212> DNA

<213> Homo sapiens

<400> 1604

<210> 1605

<211> 451

<212> DNA

<213> Homo sapiens

<400> 1605

<210> 1606

<211> 272

<212> DNA

<213> Homo sapiens

<400> 1606

```

ccggagccca cgggtggtcat ggctgccaga gcgctctgca tgctggggct ggtcctggcc 60
ttgctgtcct ccagctctgc tgaggagtac gtgggcctgt ctgcaaacca gtgtgccgtg 120
ccagccaagg acaggggtgga ctgcgggtac ccccatgtca cccccaagga gtgcaacaac 180
cggggctgct gctttgactc caggatccct ggagtgcctt ggtgtttcaa gccctgcag 240
gaagcagaat gcaccttctg aggcacctcc ag 272

```

<210> 1607

<211> 444

<212> DNA

<213> Homo sapiens

<400> 1607

```

ccaggctggg ctcaaaactcc tcacctcaac tgatccgccc accttggcct cccaaagtgc 60
tggtattata ggtgtgagcc accgtgccca aagttaagta tttttgatca agtgttttgt 120
cttttgtgca aggcatthgt ggctctgtca tagcagagga aaacaaaaca tgcctatcaa 180
atgaatcaag tccgacctct tctcatattg agcaactaga ggtctaggaa catttccct 240
acctgtcatt ctcactctggc ataccagggtg tacatactcc ttcttattct cctctgttac 300
caagatgttg gccccattgg gtttgagggtc acgaacttca caaactccaa actcttggac 360
ctcagtgttg aaggtgaggt catagcctag tgtggagaca tcattttcca gcagataaac 420
cagaccttgg tagaagtggg aatc 444

```

<210> 1608

<211> 189

<212> DNA

<213> Homo sapiens

<400> 1608

```

caaaatccaa aacttctctt gaaaagtcca gggaccgtcc aggggagatg gggaggagat 60
atggagttag tcacctgctc cagaagatgc cagcttctct ctccaggggtg cttagtggc 120
tttgcccacc cctcactccc cagggagctc tggggacagc ttctctgcac cctgtccca 180
cccacacag 189

```

<210> 1609

<211> 426

<212> DNA

<213> Homo sapiens

<400> 1609

```

cttttgttat ccttagagga ctcaactggtt tcttttcata agcaaaaagt acctcttctt 60
aaagtgcact ttgcagacgt ttcactcctt ttccaataag cttgagttag gagcttttac 120
cttgtagcag agcagtatta acacctagtt ggttcacctg gaaaacagag aggtgaccg 180
tggggctcac catgcggatg cgggtcacac ggaatgctgg agagatgtta tgtaatatgc 240
tgaggtggcg acctcagtgg agaaatgtaa agactgaatt gaattttaag ctaatgtgaa 300
atcagagaat gttgtaataa gtaaagtccct taagagtatt taaaatatgc ttccacattt 360
caaaatataa aatgtaacat gacaagagat tttgcgtttg acattgtgtc tgggaaggaa 420
gggcca 426

```

<210> 1610

<211> 447

<212> DNA

<213> Homo sapiens

<400> 1610

```

cagggctata gtgcgctatg ttgatctggt gttcatgcta agttccgcat caatatgggtg 60
acttcttggg agtggggggac caccagggttg cctaaggagg ggtgaacctg cctacgttgg 120
aaatagagct ggtcaaaact cctgtgctca tcagtagtag aattgcacct gtgaatagcc 180
accgccctcc agcatgggca acatagcaag accctgcctc ttaagataaa aattggaaaa 240
cactggtagg aaaaaaaggc tgtttggtct aaataagtct ggattgggta taaatgacac 300
aaaactatca tgaatttgaa agcatttcta atttcttgaa agtctgaaaa agtttaaaca 360
gaatttttagc tgaaaagtcc tgaaagacat ttgaaaaaaa acagcaagaa cacttaaaac 420
tattcaaggt ttgggctggg cacagtg 447

```

<210> 1611

<211> 238

<212> DNA

<213> Homo sapiens

<400> 1611

```

ccaccgggggt tgacctctct cgctagcagg gcccaccag ctcactcccc gcgtcttcca 60
tccccctctag gattccccatt gtccccctact ccagcactag gcaggcacc cagcccact 120
gcgactccca ccacgaagga cccagccct ctctcagcca acacggcccc gccaccgtc 180
tcagacatcg tgcttcttct ggtggggccag gagtctctcc tcgtcgtcga aggtctgg 238

```

<210> 1612

<211> 293

<212> DNA

<213> Homo sapiens

<400> 1612

```

ctgctgcttg tatcctcggg agaggggttc ccactctgag cgggtgggaa ggcaatgcca 60
aacatccggg aaaaataaaa ccactgtctc cacatgagct ggaactgtac gcccttgtg 120
ggtctcctca gggcgatggt agcgaatctc tgcaaaacgg taccattgtg tgcacacact 180
tagatcaatg cctgtcagag cttacaaca acgaatagca gtcttaatca acacagaggg 240
atctttttct gggctctggtc catccaacga aggagaccag tggcccccaa tgg 293

```

<210> 1613

<211> 224

<212> DNA

<213> Homo sapiens

<400> 1613

```

ctggattgac cccaaccaag gctgcaacct ggatgccatc aaagtcttct gcaacatgga 60
gactggtgag acctgcgtgt accccactca gccagtggtg gccagaaga actggtacat 120
cagcaagaac cccaaggaca agaggcatgt ctggttcggc gagagcatga ccgatggatt 180
ccagttcgag tatggcggcc agggctccga ctctgccgat gtgg 224

```

<210> 1614

<211> 439

<212> DNA

<213> Homo sapiens

<400> 1614

```

ctccaccctg gcgatggctc cctggtccta ctttctctct caaactggct ttttctcatt 60
cctttgactc cgccagactt cctcgcccc atgacctggt gttgtgtctg atcacccaa 120
cattcctggc tgcccaatgt ggggcaatga agacccagtg gaaggaatgc tagagtgtgt 180
gaaagtggag gacgcatcgt caaaggacac ctgaggacgt ctcaaagaag ctcggcggga 240
gagctgagcg ctcggaagaa ccaagaatca tctcttttga aaaatcgatt catcaaatga 300

```

```

atcttcggcc aacaactggt caagaaggat tcaaatatca caggttccaa gaagtaaagc 360
tttgagggtc acaaaattag caatagaagc tgggttcgcg catatagatt ctgctcattt 420
atacaaataa tgaggagca                                     439

```

<210> 1615

<211> 237

<212> DNA

<213> Homo sapiens

<400> 1615

```

aggcactcct ggaagtgggt cagtcagggt gcaaaaacat tgaacttgct gtcatgaggc 60
gagatcaatc cctcaagatt ttaaatacctg aagaaattga gaagtatggt gctgaaattg 120
aaaaagaaaa agaagaaaac gaaaagaaga aacaaaagaa agcatcatga tgaataaaat 180
gtctttgctt gtaattttta aattcatatc aatcatggat gagtctcgat gtgtagg    237

```

<210> 1616

<211> 266

<212> DNA

<213> Homo sapiens

<400> 1616

```

ctgggctcta gtttcattcc atctgtcatt ctcaggtaac agggacacat gtccaagtgt 60
tggtcccccgt ggcattgatt tagctttggt gataggcatt gcatcttttg tgtaatatgc 120
aataatggca tgaccagatt catgatatgc tgtgatgggt ttgtttttgt tatcaatttc 180
cacacttctt ctttcaggcc ccattagaat tttgtctttg gaaaactcca gtccttcat 240
ggttaaccatt tcttttccat caacag                                     266

```

<210> 1617

<211> 185

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (1)...(185)

<223> n = A,T,C or G

<400> 1617

```

ccatggctag gtttatagat agttgggtgg ttggtgtaaa tgagtgaggc aggagtcga 60
gnaggtagt tgtggcaata aaaatgatta aggatactag tataagagat caggttcgtc 120
cttttagtggt gtgtatgggt atcattttgt ttgagggttag tttgattagt cattgttggg 180
tggtg                                             185

```

<210> 1618

<211> 354

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (1)...(354)

<223> n = A,T,C or G

<400> 1618

```

ctgttaacag ataagtttaa cttgcatctg cagtattgca tgttagggat aagtgcttat 60
ttttaagagc tgtggagttc ttaaatatca accatggcac tttctcctga ccccttcctt 120
aggggatttc aggattgaga aatttttcca tcgagccttt ttaaaattgt aggacttggt 180
cctgtgggct tcagtgatgg ngatagtaca catntcactc agagngcatn tntgcatctt 240
ntaanatana tttcttaaaa gcctctaaag tgatcagntg ccttgatgcc aactaaggaa 300
atttgtttag cattgaatct ctgaaggctc tatgaaagga atagcatgat gtgc 354

```

<210> 1619

<211> 170

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (1)...(170)

<223> n = A,T,C or G

<400> 1619

```

ctgtgctgtg gagagaagct gatgttttgg tgtattgtca gccatcgtec tgggactcgg 60
agactatggc ctgcgcctccc caccctcctc ttggaattac aagccctggg gtttgaagct 120
gactttatag ctgcaagtgt atctnncttt tatctggtgc ctctcaaac 170

```

<210> 1620

<211> 386

<212> DNA

<213> Homo sapiens

<400> 1620

```

cctgttgatt gcatactgta gaagatttga tgttcagact gggtcttctt acatatacta 60
tgtttcgtct acagttggta aatttttgtt tttctttgta ttaaattgtg aattgtattg 120
tctggaggaa aagacagagg tctaaaaata aagaaggagt acagtttggg catggtgggt 180
cacccttgga gtcctagcac tttggggggc aaggcaggca gattgcttga gcccaggagt 240
tctagatgag cctgggcaac atagtggagc cccatctcta aaaaaacagt tttagggcca 300
ggcacagtgg ctcacacctg taagcccagc actttgggag gccgaggcag gcagatcata 360
agggcaagag attgagacca tcctgg 386

```

<210> 1621

<211> 346

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (1)...(346)

<223> n = A,T,C or G

<400> 1621

```

ccaattctgc cgttcccccg tgggccaaca acaactggggt tgtatgcgtc tggaaccctg 60
tgatagtctt cggttgcca gcctggccca ccacatccac tgcttgccc acacggacag 120
acactggcaa tggccgcagc tcctcatcaa acgtaaccag cattcggggc tgcattggcag 180
ccaccagccc atacaatata tagtgtgatt tgcctagaat aatgtttcga acatccagga 240
aagagacaag cacagtgagc agtccancca cggccacctg gtcataagc tgccgggtcgc 300
tgtggtaggg gcagagggta aggggtgcct tccctaaatg tgtcag 346

```

<210> 1622
 <211> 366
 <212> DNA
 <213> Homo sapiens

<400> 1622
 ggaagtttgt gctctctgcg tggctaagtt tttcacctac taggacgggg gtgggggtggg 60
 gagaacaggt gtccttctaa aatacagcac aagctacagc ctgctgccag ccataaccca 120
 ggagtaacat cagaaacagg tgagaatgac cactttaact caccggggccc gtcgcactga 180
 aataagcaag aactctgaaa agaagatgga aagtgaggaa gacagtaatt gggagaaaag 240
 tccagacaat gaagattctg gagactctaa ggatatccgc cttactctta tggaagaagt 300
 attgcttctg ggactaaaag ataaagaggg gtacacatct ttctggaatg actgcatatc 360
 atcagg 366

<210> 1623
 <211> 165
 <212> DNA
 <213> Homo sapiens

<400> 1623
 ctgttgattg gctgtgacac tgctttgtgt catcttctta ccatgatcaa aggcgaagga 60
 agggatctct tttgggacat tgtgattgtt ttagcagaga gagaaagaga tgaaatacac 120
 ttcggttttc tcttaaaaga tgcattgtatc atacagtgtc ttaag 165

<210> 1624
 <211> 227
 <212> DNA
 <213> Homo sapiens

<400> 1624
 ccaatgcccc gagcaggccc tctttccatc cctgtcggga tgagctgggc aactatgtca 60
 acaaacggaa taccacgtgg caagccgggc acaacttcta caacgtggac atgagctact 120
 tgaagaggct atgtggtacc ttcctgggtg ggcccaagcc accccagaga gttatgttta 180
 ccgaggacct gaagctgcct gcaagcttcg atgcacggga acaatgg 227

<210> 1625
 <211> 373
 <212> DNA
 <213> Homo sapiens

<400> 1625
 ctgtagcttt tgtgggactt ccaactgctca ggcgtcaggc tcaggtagct gctggccgcg 60
 tacttgttgt tgctttgttt ggagggtgtg gtggtctcca ctcccgcctt gacggggctg 120
 ctatctgcct tccaggccac tgtcacggct cccgggtaga agtcacttat gagacacacc 180
 agtgtggcct tgttgcttg aagctcctca gaggaggggtg ggaacagagt gaccgagggg 240
 gcagccttgg gctgacctag gacggctcagt ttggtccctc cgccgaacac ccgaagataa 300
 ttagtgctgt ctgttgagta acaatagtag tcaccttcac cttccacctg ggccccagtg 360
 atgggtcaagg tgg 373

<210> 1626
 <211> 367
 <212> DNA
 <213> Homo sapiens

<400> 1626

```
ccagacgtgg tggctcacac ctgcaatccc agcaccttag gaggccgagg caggaggatc 60
cttgaggtca ggagttcgag accagcctcg ccaacatggg gaaaccccat ttctactaaa 120
aatacaaaaa ttagccaagt gtggtggcat atgcctgtaa tcccaactac tcagaaggcc 180
gaggcaggag aattacttga acgcaggaga atcactgcag ccctggaggc agaggttgca 240
gtgagccgag attgcaccac tgtactccag cctgggtgac agagcaagac tccatctcag 300
taaataaata aataaataaa aagcgctgca gtagctgtgg cctcaccctg aagtcagcgg 360
gcccagg                                           367
```

<210> 1627

<211> 424

<212> DNA

<213> Homo sapiens

<400> 1627

```
ctggataagg acatcaatac cttctctatg cgtgtcaggg tgtgggtacgg gtatcacttt 60
ccggagctgg tgaagatcat caacgacaat gccacatact gccgtcttgc ccagttttatt 120
ggaaaccgaa ggaactgaa tgaggacaag ctggagaagc tggaggagct gacaatggat 180
ggggccaagg ctaaggctat tctggatgcc tcacggctct ccattgggcat ggacatatct 240
gccattgact tgataaacat cgagagcttc tccagtcgtg tgggtgtcttt atctgaatac 300
cgccagagcc tacacactta cctgcgctcc aagatgagcc aagtagcccc cagcctgtca 360
gccctaattg gggaagcggg aggtgcacgt ctcacgcac atgctggcag cctcaccaac 420
ctgg                                           424
```

<210> 1628

<211> 314

<212> DNA

<213> Homo sapiens

<400> 1628

```
tcgactgtta tagcttagaa agcaacacta ctactatgag actataaaac attaaactat 60
tttaagaaaa ccacgtgtg gaaaaatgga gccatttttg tcaaaaagtg gctcaaagca 120
caaaactgct cagatgttca agagtcctag gagtctgggc tgcacagtat taaggggtga 180
gaggagaccg acagcctgtt tgaatcaggc ttgtgagccc agctcatctg acaacttcaa 240
agagcttctc tgcctataca ttccaccgtt tagcataaga caccacttta cgctattttac 300
aagtctcctt ttgg                                           314
```

<210> 1629

<211> 393

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (1)...(393)

<223> n = A,T,C or G

<400> 1629

```
ctggaccagc accccattga cgggtacctc tcccacaccg agctggctcc actgcgtgct 60
cccctcatcc ccattggagc ttgcaccacc cgcttttttc agacctgtga cctggacaat 120
gacaagtaca tcgccttgga tgagtgggcc ggctgcttcg gcacaaagca gaaggatatc 180
gacaaggatc ttgtgatcta aatccactcc ttccacagta ccggattctc tctttaaccc 240
tccccttcgt gttttccccc aatgtttaaa atgtttggat ggtntgttgt tctgcctgga 300
gacaaagggt ctaacataga tttaagttga ataacattaa cgggtgctaaa aaatgaaaaa 360
```

ttctaaccga agacatgaca ttcttagctg taa

393

<210> 1630

<211> 317

<212> DNA

<213> Homo sapiens

<400> 1630

```
ctgcaagaat atcagaaatc aatacaaaca agtattgaca ggtgttacag acatgcaaaa 60
tatccttcaa tgcaacgaat ttttaagaaa tcagctagcc tatattaatc agatgtttta 120
ggtcaaacca agtttccatc tcgggctcag tgaaatagta ttaactcatt gagtctcctt 180
tccccagga atgttgggaa tggcagaaca gaaagagcta tcaactcctta aattctttta 240
tgcgagtgtt actccaacac ttattttact tggtttactt ggaatgtatg agaggaaact 300
gatgtttttt acaatgg                                     317
```

<210> 1631

<211> 262

<212> DNA

<213> Homo sapiens

<400> 1631

```
ccttaggcaa gtcaccttac ttatctaaga ctgtttcccc acctggaaga tgccctacaa 60
gcctcctgtg gctgtgttta gaaagcatgc ccggcctttc ttgacagcca gccaccccag 120
atgatggcag ggcaagggaag actgttagga gtcagagtgc tccccctcagg tggaaggaaa 180
ctgggccaac tctactttgt aagccatagg gtgccaggta gcccggccac cctgagcctg 240
tgctccact gccccgcgt gg                                     262
```

<210> 1632

<211> 138

<212> DNA

<213> Homo sapiens

<400> 1632

```
ctggaattaa ttcttcgaca actccagacc gaccttcgga agggaaaaaca agacaaggcc 60
gttctccaag cagaagtgca gcacctgaga caggacaaca tgagactgca ggaggagtcc 120
cagaccgcga cagctcag                                     138
```

<210> 1633

<211> 192

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (1)...(192)

<223> n = A,T,C or G

<400> 1633

```
ccttgaaggg acctcanagc aaaggaagag acctgggtgt ggtgaggcat cccanggcac 60
ggaagggacc ggttgtgctn ngggaatcca ctgnncctc cttggnnaaa aaagcacaac 120
acatcatata tatttaccag accagaagcg ctggcccca gtctcccca cctggtcggg 180
ggaacctcct gg                                     192
```

<210> 1634

<211> 447

<212> DNA

<213> Homo sapiens

<400> 1634

```
ctgcttttaa aggtcttaaa tcaactgaat accttgactt gagcttcaat cagatagcca 60
gactgccttc tggctcctct gtctctcttc taactctcta cttagacaac aataagatca 120
gcaacatccc tgatgagtat ttcaagcggt ttaatgcatt gcagtatctg cgtttatctc 180
acaacgaact ggctgatagt ggaatacctg gaaattcttt caatgtgtca tccctgggtg 240
agctggatct gtcctataac aagcttaaaa acataccaac tgtcaatgaa aaccttgaaa 300
actattacct ggaggtcaat caacttgaga agtttgacat aaagagcttc tgcaagatcc 360
tgggggcatt atcctactcc aagatcaagc atttgcggtt ggatggcaat cgcactctcag 420
aaaccagtct tccaccgat atgtatg 447
```

<210> 1635

<211> 364

<212> DNA

<213> Homo sapiens

<400> 1635

```
gtttttatttg agacataaaa acacatgtgt ttctattaca tagtgtgggg tttaggggcc 60
tggtttctaa gacaagactt tatttcaccc tgtatcacag cttcctggga aatgaattag 120
ggagcaagag acggcctggc aagaaaatca ttattgttgc tgggaagttg caaagaaagg 180
ggagagttta ttcaaattag tgtaacagag cccccaggat gaagagagtg gtgcagggaa 240
aaggtctaaa ttcttggtgt tgggtggggac actggcacat cccacagcaa ggactcagcc 300
ctcaacggcg gcggctgggt cttgggaggg gagtgggtggg agggtaaggg ctctcagct 360
ccct 364
```

<210> 1636

<211> 399

<212> DNA

<213> Homo sapiens

<400> 1636

```
ctggctggct agactgtttg tgcgccaaga ggatggtcag cgctgctttc cagcctggct 60
ctgctggggc gctggcatct ggttcagttc caccattctc cctgctttct ttgccaaagt 120
tgatattcac ccaagggcac cagtctctat gctgagaggt gggatcaaag aagcttcggg 180
aagatgtgtc cgaactgctg gaggagcaga ggcgagctcg cttggctttc cgcagagggc 240
tagatggtac ctccaggcca ggggtgtctc ctgttcccat gcttcgggtc actgggagag 300
ttctgggtgt ggggctagca gcctctggct caggacgggt aacaggactg gaagagtccc 360
agctccgagt tcgagagaca atgggaccag ggctctttt 399
```

<210> 1637

<211> 246

<212> DNA

<213> Homo sapiens

<400> 1637

```
ctgagctttc agcagataaa tcacagcaga aatagaatca ccctaggact ttcaatcaaa 60
agctggaagt ccaccttaca gaaagacaaa aagaaacccc tttttatata ttaacaaagc 120
aatagctctc aagcagcaga gcatctcgag gaagaaagct tgcccggctg ccatcccatc 180
atgccagagc gtgcagtgtc cacccttgac tacgctgggg aattgctgat tttttgaaaa 240
agcttg 246
```

<210> 1638
 <211> 453
 <212> DNA
 <213> Homo sapiens

<400> 1638
 ccaagagttc tccactgtga agactgaaag gacctggtga catttcggca tcagtcctgt 60
 taccacttgg aggtaacaga agcaggctcg tgctctcctt taattctacc acactacatg 120
 actcgcaatt ggttctgaaa ttagaacgtt caccatcgta cttaaaatct taggggcatg 180
 aagagtcagc tagaacaagg aaaaagaaag tcgcaggtag taggtaagta ggtgggcaca 240
 tgaaaagcca agctgctctg tccaacacca gtgtacatgt gctttaacta aatgaactcc 300
 agaggccaac agcagcagac ctgctcaatt caccttccaa atcagaacaa gacaaaaag 360
 ctcaggcttg agttgtcaac tatgcatagg ttccgccagt gatgaggagc tcgtaagcag 420
 gatctctact ccttctgcac aacacgatgc aag 453

<210> 1639
 <211> 197
 <212> DNA
 <213> Homo sapiens

<400> 1639
 tttgctgttc gtgatatgag acagacagtt gcggtgggtg tcatcaaagc agtggacaag 60
 aaggctgctg gagctggcaa ggtcaccaag tctgcccaga aagctcagaa ggctaaatga 120
 atattatccc taatacctgc caccctcctc ttaatcagtg gtggaagaac ggtctcagaa 180
 ctgtttgttt caattgg 197

<210> 1640
 <211> 278
 <212> DNA
 <213> Homo sapiens

<400> 1640
 ccagagcggg gagtcccacc acctcgaact ctgggaattc gagccacagc tctgccagta 60
 cccaagact cagcactagt ctgatgacct gctaattcac tgacagcata gggctgtctg 120
 ttgtttttgc gcaagttggt gtgaacaaag ttcacaatat ctggtcgaat aggagccttg 180
 aatacagcag gcaaagtgc atttttgcca gatgactccc ccttttcgga gtacaccgat 240
 atcagtgggc gagcgcacgc catggcggac ctcggccg 278

<210> 1641
 <211> 227
 <212> DNA
 <213> Homo sapiens

<400> 1641
 ccattgttcc cgtgcatcga agcttgacag cagcttcagg tcctcggtaa acataactct 60
 ctgggggtggc ttgggccac ccaggaagg accacatagc ctcttcaagt agctcatgtc 120
 cacgtttaga aagttgtgcc cggcttgcca cgtggtattc cgtttgttga catagttgac 180
 cagctcatcc gacaggggat ggaaagaggg cctgctccgg gcattgg 227

<210> 1642
 <211> 299
 <212> DNA
 <213> Homo sapiens

<400> 1646

<210> 1647

<212> DNA

<213> Homo sapiens

<400> 1647

ccagcttgca	agcacgctgg	caaattctctg	tcagggtcagc	tccagagaag	ccatttagtca	60
tttttagccag	gaactccaag	tccacatcct	tggcaactgg	ggacttgcgc	aggttagcct	120
tgaggatggc	aacacgggac	ttctcatcag	gaagtgggat	gtagatgagc	tgatcaagac	180
ggccagggtct	gaggatggca	ggatcaatga	tgtcaggccg	gttggtagcg	ccaatgatga	240
acacatTTTT	ttttgtggac	atgccatcca	tttctgtcag	gatctggttg	atgactcggt	300
cagcagcccc	accaccatct	ccaatgttac	ctccacgagc	cttggcaatc	gaatccagct	360
catcaaaagaa	tagcacacag	ggggcagctt	gcggggcctt	gtcaaagatt	tctctgacat	420
tggcctcaga	ctccccaaac	ccacatggtga	g			451

<210> 1648

<211> 176

<212> DNA

<213> Homo sapiens

<400> 1648

cctaaacgag gatttcagct tccattatgc ccaactccag tccaacatca ttgaggcgat 60
taatgagctg ctagtggagc tggaagggaac aatggagaac attgcagccc aggcctctgga 120
gcacattcac tccaatgaqg tqatcatgac cattggcttc tcccgaaacag taqaqg 176

<210> 1649

<211> 435

<212> DNA

<213> Homo sapiens

<400> 1649

tgtggtgtg	cgttggtcc	tgtgcggtca	cttagccaag	atgcctgagg	aaaccagac	60
ccaagaccaa	ccgatggagg	aggaggaggt	tgagacgttc	gcctttcagg	cagaaattgc	120
ccagttgatg	tcattgatca	tcaatacttt	ctactcgaac	aaagagatct	ttctgagaga	180
gctcatttca	aattcatcag	atgcattgga	caaaatccgg	tatgaaaagt	tgacagaccc	240
cagtaaatta	gactctggga	aagagctgca	tattaacctt	ataccgaaca	aacaagatcg	300
aactctcact	attgtggata	ctggaattgg	aatgaccaag	gctgacttga	tcaataacct	360
tggtactatc	gccaaagtctg	ggaccaaagc	gttcatggaa	gctttgcagg	ctggtgcaga	420
tatctctatg	attgg					435

<210> 1650

<211> 246

<212> DNA

<213> Homo sapiens

<400> 1650

```
ccatgtctgt attgtaactg gtaaaaggct tcaagtcaga ttgatgatca agaaaagtca 60
aaaccccagc ccaagattgg gaaagcaggt ggtggttcca agctttttaa aaattattga 120
agctctccat cctgttctgt gagtgtgtct tctctttctc cttcacgtca tagccgtgac 180
ccaccgttca tctctgtctt tgcgtaaaga tgaccgatgg agtccaaagc caagtggctt 240
caccag                                           246
```

<210> 1651

<211> 400

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (1)...(400)

<223> n = A,T,C or G

<400> 1651

```
cggcaagttc tcccaggaga aagccatgtt cagttcgagc gccaaagaccg tgaagcccaa 60
tggcgagaag ccggacgagt tcgagtccgg catctcccag gctcttctgg agctggagat 120
gaactcggac ctcaaggctc agctcaggga gctgaatatt acggcagcta nngaaattga 180
agttggtggt ggtcggaaaag ctatcataat ctttgttccc gtctctcaac tgaaatcttt 240
ccagaaaatc caagtccggc tagtacgcga attggagaaa aagttcagtg ggaagcatgt 300
cgnctttatc ggctcagagg aggaattctg cctaagccaa ctcnaaaaag ccgnacnaaa 360
aattanngca aaaagcgtnc caggagccgt nctctgacag                                           400
```

<210> 1652

<211> 338

<212> DNA

<213> Homo sapiens

<400> 1652

```
ctgggggtgc ccattcttctg tgctctgtgg tacatatctg tgtcgccaaa gtagcgtgcc 60
cggtagacga agccttctct ctgctgcttc tcttccagc agttgttccg gaggttggcg 120
atataatcat cttccacatt ccgctcgact gttttgaggc tggagcctgt gtactcttcg 180
gagaaagtgt ctccacata gtagacgaca cccaggtggt cagtgaactg cctgtggatg 240
tggccacagc acggtcttgg actcagactg tagggtggac tggagaccat gagctggctg 300
agagctgaca cgagaatcag gatgaggata ggcacacag                                           338
```

<210> 1653

<211> 167

<212> DNA

<213> Homo sapiens

<400> 1653

```
gcggtggagc cgccaccaa atgcagattt tcgtggaaac ccttacgggg aagaccatca 60
ccctcgaggt tgaaccctcg gatacgatag aaaatgtaaa ggccaagatc caggataagg 120
aaggaattcc tcttgatcgg cagagactga tctttgctgg caagcag                                           167
```

<210> 1654
 <211> 1034
 <212> DNA
 <213> Homo sapiens

<220>
 <221> misc_feature
 <222> (1)...(1034)
 <223> n = A,T,C or G

<400> 1654
 atgcatgctc gagcgggccgc cagtgtgatg gatatactgca gaattcgccc ttagcgtggg 60
 cgcgggccgag gtccaagagg gagataaanac aaactttctca aacaaaaaga aaagaaaaac 120
 gaatgattca tctgctttta tcagtgtgat taatgcagca cccattgccc cggaaccgt 180
 ttctgctgta ctatctggat actaaaatgt tacggaagta gctctttgtt ctccctcact 240
 ctgcccttag ttaatagaaa ttcagactcg ccaagtaagg ctttgtgcat agtgtcttca 300
 tgtcgcgat agttgagcgc gttcttagca gttggcttca tggacagctc attagtgttt 360
 tgacttttct taccagcgt taattgaatt cttgctttta gacaacttcc tttttgtagt 420
 ggtgaacctt gcccttttagt acagttcaag tgaatctgga taattgttca tctttgcttt 480
 agcttagata ccatgtagtg gtctgtggct acaggaagct gggtctgtct gcttccacag 540
 tctgcttaaa aaactgtctg acttcgtgaa tatagagacc aagtttacca cttctgatga 600
 agagaccaat taagattcat tcctcattct gtttctttcc agtgggagaa gagtccccat 660
 gaaataagat gaaactgatt ccatgcacta gtacatgtag gcttctccct tgcgcaaagc 720
 ttaacaattt gtaggaaact ttgggtcttt ttgtcccaag aaaaaggaat gtcttgacag 780
 gcttaaagct ttctgtcccc ttgcacctta aaactcgaaa gttagnaaa atccctttaa 840
 agggcttttt ttaatagcca gaacttccca aaaggaatgg cnttttaggg aatttcntag 900
 ccatngcttt ttaaatttaa agaaattttt aanaaccttg cccnnggggn ggggncccg 960
 tccaaaaagg gnggnaaaaa ttcccagcc naccctttng gggggggccn cgttttccct 1020
 tnnngggggg aanc 1034

<210> 1655
 <211> 487
 <212> DNA
 <213> Homo sapiens

<400> 1655
 atgcatgctc gagcgggccgc cagtgtgatg gatatactgca gaattcgccc tttcgagcgg 60
 ccgcccgggc aggtcctact cttctccgctc cattgtacta tctgcccgtg gtggggatgg 120
 cagtaggac atatttgatg acttccgaga agcatattat tggctccgctc ataatactcc 180
 agaggatgag aaggtcatgt cctggtggga ttatggctat cagattacag ctatggcaaa 240
 ccgaacaatt ttagtgagca ataacacatg gaataatacc catatttctc gagtagggca 300
 ggcaatggcg tccacagagg aaaaagccta tgagatcatg agggagctcg atgtcagcta 360
 tgtgctgggc atttttgag gacctcggcc gcgaccacgc taaggcgaa ttccagcaca 420
 ctggcgccg ttactagtgg atccgagctc ggtaccaagc ttggcgtaat catggtcata 480
 gctgttt 487

<210> 1656
 <211> 514
 <212> DNA
 <213> Homo sapiens

<220>
 <221> misc_feature
 <222> (1)...(514)

<223> n = A,T,C or G

<400> 1656

```
atgcatgctc gagcgggccc ccagtgtgat ggatatctgc agaattcgcc cttancgtgg 60
tcgcggccga ggtcctaccc ataatccaga gaggcttgcc cagaggagga ctacgtgggg 120
gacgtgccac cagaacccta cttgggggcg ggatgtcact ccgaggtcaa aacctgctcc 180
gaggtggacg agccgtagct ccccgaatgg gcttaagaag aggtggtggt cgaggtcgtg 240
gaggtcctgg gagagggggc ctagggcgtg gagctatggg tcgtggcgga atcgggtggt 300
gaggtcgggg tatgataggt cggggaagag ggggctttgg aggccgaggc cgaggccgtg 360
gacgagggag aggtgccctt gctcgccctg tattgaccaa ggagcagacc tgcccggggc 420
gccgctcgaa gggcgaattc cagcacactg gcggccgtta ctagtggatc cgagctcggg 480
accaagcttg gcgtaatcat ggtcatagct gttt 514
```

<210> 1657

<211> 605

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (1)...(605)

<223> n = A,T,C or G

<400> 1657

```
atgcatgctc gagcgggccgc cagtgtgatg gatattctgca gaattcgccc tttcgagcgg 60
ccgcccgggg aggtccanac gctgacattg nttctgagtc cttaagcagg aaggatttga 120
aatcctggag cttggcagtc ttgctcttca cctctaagcc aatggtgacc ccttcattca 180
taaagtcac aactctccgg aagtcaccc caccgaactg tcgagaagtt aaggctgggg 240
ccccaagccg caggccgccc ggtgtgatgg cacttcggtc tccaggacag gtgttcttgt 300
tggcagtgat ggatacaagc tctagcaccg gctcagcccg agctccatcc aggcccttgg 360
gccgcaggtc caccagcacc aggtggttgt cagtaccacc tgataccagt gagtgcctc 420
gccctagcag ggcattctgcc atggcccag cattcttcag aacctgcagg gagtactccc 480
ggaacatggg ggtgcaggac ctcgcccgcg accacgctaa gggcgaattc cagcacactg 540
gcggccgtta ctagtggatc cgagctcggg accaagcttg gcgtaatcat ggtcatagct 600
gtttc 605
```

<210> 1658

<211> 784

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (1)...(784)

<223> n = A,T,C or G

<400> 1658

```
agnnttcn cggccctcna gntgcatgct cgagcggccg cgcagtgaga tgnatatctg 60
cagaattcgc cttancgtg ggcgnangca tgacgctcgg gatcagaact aaaacaagt 120
agatcacccc tctaattatt tctgaactng gttaataaaa gcttataaga tttttatgaa 180
gcancactg tatgatattt taagcaaata tggtatttaa aatattgatc cttcccttgg 240
accaccttca tgtagttgg gtattataaa taagagatac aacctgaat atattatgtt 300
tatacaaaat caatctgaac acaattcata aagatttctc ttttatacct tcctcactgg 360
ccccctccac ctgcccatag tcaccaaatt ctgttttaaa tcaatgacct aagatcaaca 420
```


tcctnggcat agctgtttc

559

<210> 1661

<211> 453

<212> DNA

<213> Homo sapiens

<400> 1661

```

ttgggccctc tagatgcatg ctcgagcggc cgccagtgtg atggatatct gcagaattcg 60
cccttttcgag cggccgcccc ggccaggtctg cagtgtccct ttttatatca tgctagtgtt 120
gagacatact tgactaactt gggaacagtt cgatatattg acaaccgtca acttaagaaa 180
atcaacagct tttggcccca gcgtccaagt gaacttttca tggagtgcag aatctcaaat 240
ggacaaaata ctttgtcttt ttaaatactg aaaatttaat tattagtact atgactgaaa 300
gattcttcat ggctaaaaag ctctgcatca aactcaattc aggaggacct cggccgcgac 360
cacgctaagg gcgaattcca gcacactggc ggccgttact agtggatccg agctcggtag 420
caagcttggc gtaatcatgg tcatagctgt ttc 453

```

<210> 1662

<211> 809

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (1)...(809)

<223> n = A,T,C or G

<400> 1662

```

ctcgagcggc cgccantgtg atggntatct gcagaattcg cccttanccg ccgcccgggc 60
aggtecttag ccaaagaatg cagtggagcc ttccccnngg ggctgcattg tgaatgaata 120
ccaattgaca gcataaaaat taatagtccc atatcagatc tggaaggggt ttctggggct 180
gtctgatgtc cctatcctgt ttagtggaac acaatagcag aaaattcttt ctgggtccat 240
ctgctataaa gtcttggtta aacagcatta ctatgaagag gatgaactca cctaccttca 300
natggaggaa aagtgaagag gacttaggct ttagtctctc atgacttttc ttaagcacta 360
cctacctgta ataagctgag tgcaaaagga tgccgaagaa aatctgcacc cagaagctgt 420
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agattctttg ntcaaggggn actttgccaa aaggggcaga atangngggg aaagagttgc 540
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ggntataatt aaaatgnngc tttttacact gnggggggcta tataaaaact gggtagnaaa 660
atttccaccg agcatntatg gngatttgnt cacagnaaac ctccgggcng gaccacgct 720
aagggnggaa ttccagcnac antggggggg ncngntacct anagtggatc ccnagnctng 780
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<210> 1663

<211> 585

<212> DNA

<213> Homo sapiens

<400> 1663

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gatattctaca aggctaataa cattgcctat gaagatgtgg tcggggggaga agactggaac 180
ccagtagagg agaaaataga gagtcaaacc caggaagagg tgagagacag caaagagaat 240
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ggggaaaggg ccaccaggct ttttgagaaa cctcttgatt ctcagtttat ttatcagacc 480
tcggccgcga ccacgctaag gggaattcc agcacactgg cggccgttac tagtggatcc 540
gagctcggta ccaagcttgg cgtaatcatg gtcatactg tttcc 585

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<210> 1664

<211> 999

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<222> (1)...(999)

<223> n = A,T,C or G

<400> 1664

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agtaagaaca cactaacgtc acaagtttct cattctaaag tgcnaaancc ntaatngtct 540
ngaaagtgga acaggggtaa agggcaaaaa ttaaccccc ccacccaat taaagtttcc 600
tggaangtca ntantntttt naatcccaa aggnnncatt tctntttaaa aaaattggnt 660
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cacccttgng gaaacncttt tngtgggggn cccggtcgna aaacccaacc nccctntaaa 840
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cgntttaccc nttaaaatgg gggaattcc ccgaaagcgt ttgggggtaa ccccaaaaga 960
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<210> 1665

<211> 27

<212> DNA

<213> Artificial Sequence

<220>

<223> PCR primer

<400> 1665

gctaaagggtg accccaagaa accaaag

27

<210> 1666

<211> 37

<212> DNA

<213> Artificial Sequence

<220>

<223> PCR primer

<400> 1666

ctattaactc gagggagaca gataaacagt ttccttta

37

<210> 1667

<211> 207

<212> PRT

<213> Homo sapiens

<400> 1667

Met	Gln	His	His	His	His	His	His	Ala	Lys	Gly	Asp	Pro	Lys	Lys	Pro
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Lys	Gly	Lys	Met	Ser	Ala	Tyr	Ala	Phe	Phe	Val	Gln	Thr	Cys	Arg	Glu
			20					25					30		
Glu	His	Lys	Lys	Lys	Asn	Pro	Glu	Val	Pro	Val	Asn	Phe	Ala	Glu	Phe
			35				40					45			
Ser	Lys	Lys	Cys	Ser	Glu	Arg	Trp	Lys	Thr	Met	Ser	Gly	Lys	Glu	Lys
			50			55					60				
Ser	Lys	Phe	Asp	Glu	Met	Ala	Lys	Ala	Asp	Lys	Val	Arg	Tyr	Asp	Arg
65					70				75					80	
Glu	Met	Lys	Asp	Tyr	Gly	Pro	Ala	Lys	Gly	Gly	Lys	Lys	Lys	Lys	Asp
				85					90					95	
Pro	Asn	Ala	Pro	Lys	Arg	Pro	Pro	Ser	Gly	Phe	Phe	Leu	Phe	Cys	Ser
			100					105					110		
Glu	Phe	Arg	Pro	Lys	Ile	Lys	Ser	Thr	Asn	Pro	Gly	Ile	Ser	Ile	Gly
			115			120						125			
Asp	Val	Ala	Lys	Lys	Leu	Gly	Glu	Met	Trp	Asn	Asn	Leu	Asn	Asp	Ser
			130			135					140				
Glu	Lys	Gln	Pro	Tyr	Ile	Thr	Lys	Ala	Ala	Lys	Leu	Lys	Glu	Lys	Tyr
145					150					155				160	
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				165					170					175	
Lys	Gly	Pro	Ala	Lys	Val	Ala	Arg	Lys	Lys	Val	Glu	Glu	Glu	Asp	Glu
			180					185					190		
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<210> 1668

<211> 636

<212> DNA

<213> Homo sapiens

<400> 1668

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tccgggaaag	agaaatctaa	atttgatgaa	atggcaaaag	cagataaagt	gcgctatgat	240
cgggaaatga	aggattatgg	accagctaag	ggaggcaaga	agaagaagga	tcctaagtct	300
cccaaaaggc	caccgtctgg	attcttctctg	ttctgttcag	aattccgccc	caagatcaaa	360
tccacaaacc	ccggcatctc	tattggagac	gtggcaaaaa	agctgggtga	gatgtggaat	420
aatttaaagt	acagtgaaaa	gcagccttac	atcactaagg	cggcaaagct	gaaggagaag	480
tatgagaagg	atgttgctga	ctataagtgc	aaaggaaagt	ttgatggtgc	aaaggggtcca	540

gctaaagttg cccggaaaaa ggtggaagag gaagatgaag aagaggagga ggaagaagag 600
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<210> 1669

<211> 2821

<212> DNA

<213> Homo sapiens

<400> 1669

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<210> 1670
 <211> 137
 <212> PRT
 <213> Homo sapiens

<400> 1670
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 Gly Ala Pro Glu Gly Pro Gly Pro Ser Gly Gly Ala Gln Gly Gly Ser
 20 25 30
 Ile His Ser Gly Arg Ile Ala Ala Val His Asn Val Pro Leu Ser Val
 35 40 45
 Leu Ile Arg Pro Leu Pro Ser Val Leu Asp Pro Ala Lys Val Gln Ser
 50 55 60
 Leu Val Asp Thr Ile Arg Glu Asp Pro Asp Ser Val Pro Pro Ile Asp
 65 70 75 80
 Val Leu Trp Ile Lys Gly Ala Gln Gly Gly Asp Tyr Phe Tyr Ser Phe
 85 90 95
 Gly Gly Cys His Arg Tyr Ala Ala Tyr Gln Gln Leu Gln Arg Glu Thr
 100 105 110
 Ile Pro Ala Lys Leu Val Gln Ser Thr Leu Ser Asp Leu Arg Val Tyr
 115 120 125
 Leu Gly Ala Ser Thr Pro Asp Leu Gln
 130 135

<210> 1671
 <211> 109
 <212> PRT
 <213> Homo sapiens

<400> 1671
 Met Ala Arg Pro Glu Leu Arg Pro Gly Gly Gly Gly Glu Ser Arg Gly
 5 10 15
 Gly Gly Asp Asp Gly Ala Ala Cys Arg Arg Asn Ala Gly Gln Gly Arg
 20 25 30
 Arg Gly Ser Gly Gly Ala Arg Gly Ala Arg Ala Glu Arg Arg Arg Ala
 35 40 45

Gly Arg Gln His Pro Leu Gly Pro His Arg Arg Gly Ala Gln Arg Ala
 50 55 60

Ala Glu Arg Ala His Pro Ala Ala Ala Val Arg Val Gly Pro Arg Gln
 65 70 75 80

Gly Ala Glu Pro Arg Gly His Asp Pro Gly Gly Pro Arg Gln Arg Ala
 85 90 95

Pro His Arg Cys Pro Leu Asp Gln Arg Gly Pro Gly Arg
 100 105

<210> 1672

<211> 145

<212> PRT

<213> Homo sapiens

<400> 1672

Met Gly Leu Lys Ser His Val Leu Pro Ala Pro Asn Ser Gln Gly Gln
 5 10 15

Gly Ser Leu Cys Ile Phe Val Tyr Val Thr Ser Tyr Met Asp Tyr Ile
 20 25 30

Gln Leu Gln Gly Lys Glu Asn Leu Asp Cys Ser Gly Leu Asn Lys Gln
 35 40 45

Lys Ile Val Phe Pro His Ser Met Asp Ser Gly Asp Gly Trp Leu Met
 50 55 60

Val Leu Val Gln Gln Leu His Glu Gly Arg Gly His Val Leu Asp Pro
 65 70 75 80

Phe Ala Leu Ile Ser Val Leu Val Thr Ser Trp Ser Gln Asp Gly Cys
 85 90 95

Cys Ile Pro Lys Asn His Val Cys Val Gln Gly Arg Arg Gly Gly Gly
 100 105 110

Arg Gly Arg Ala Lys Leu Ala Gly Pro Val Thr Phe Tyr Gln Lys Val
 115 120 125

Lys Pro Arg Gln Lys Ser Val Ser Cys Ser Leu Pro Leu His Ile Phe
 130 135 140

Thr
 145

<210> 1673

<211> 117

<212> PRT

<213> Homo sapiens

<400> 1673

Met Asp Tyr Ile Gln Leu Gln Gly Lys Glu Asn Leu Asp Cys Ser Gly
 5 10 15

Leu Asn Lys Gln Lys Ile Val Phe Pro His Ser Met Asp Ser Gly Asp
 20 25 30

Gly Trp Leu Met Val Leu Val Gln Gln Leu His Glu Gly Arg Gly His
 35 40 45

Val Leu Asp Pro Phe Ala Leu Ile Ser Val Leu Val Thr Ser Trp Ser
 50 55 60

Gln Asp Gly Cys Cys Ile Pro Lys Asn His Val Cys Val Gln Gly Arg
 65 70 75 80

Arg Gly Gly Gly Arg Gly Arg Ala Lys Leu Ala Gly Pro Val Thr Phe
 85 90 95

Tyr Gln Lys Val Lys Pro Arg Gln Lys Ser Val Ser Cys Ser Leu Pro
 100 105 110

Leu His Ile Phe Thr
 115

<210> 1674

<211> 90

<212> PRT

<213> Homo sapiens

<400> 1674

Met Asp Ser Gly Asp Gly Trp Leu Met Val Leu Val Gln Gln Leu His
 5 10 15

Glu Gly Arg Gly His Val Leu Asp Pro Phe Ala Leu Ile Ser Val Leu
 20 25 30

Val Thr Ser Trp Ser Gln Asp Gly Cys Cys Ile Pro Lys Asn His Val
 35 40 45

Cys Val Gln Gly Arg Arg Gly Gly Gly Arg Gly Arg Ala Lys Leu Ala
 50 55 60

Gly Pro Val Thr Phe Tyr Gln Lys Val Lys Pro Arg Gln Lys Ser Val
 65 70 75 80

Ser Cys Ser Leu Pro Leu His Ile Phe Thr
 85 90

<210> 1675

<211> 102

<212> PRT

<213> Homo sapiens

<400> 1675

Met Gln Asn Cys Val Pro Val Ser Phe Cys Cys Val Thr Asn His Pro
 5 10 15

Gln Thr Trp Gln Leu Glu Thr Asn Pro Val Phe Ser His Asn Pro Met
 20 25 30

Gly Trp Gln Phe Gly Leu Gly Ser Thr Gly Gln Phe Cys Cys Ser His
 35 40 45

Leu Gly Ser Leu Met Glu Leu Arg Ser Ala Val Thr Ser Ala Gly Pro
 50 55 60

Gly Trp Ser Arg Ile Ala Leu Leu Thr Cys Leu Ala Gly Asp Arg Leu
 65 70 75 80

Leu Ala Gly Ile Ala Trp Phe Ser Ser Met Trp Pro Leu Gln Gln Ala
 85 90 95

Ser Ser Gly Leu Phe Thr
 100

<210> 1676

<211> 1336

<212> DNA

<213> Homo sapiens

<400> 1676

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 tctacggagg ctttcccttc tggggaggct tgtggtttat catttcagga tctctctccg 540
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<210> 1677
<211> 250
<212> PRT
<213> Homo sapiens
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Val Ala Pro His Asn Gly Tyr Pro Val Thr Pro Gly Ile Met Ser His
20 25 30

Val Pro Leu Tyr Pro Asn Ser Gln Pro Gln Val His Leu Val Pro Gly
35 40 45

Asn Pro Pro Ser Leu Val Ser Asn Val Asn Gly Gln Pro Val Gln Lys
50 55 60

Ala Leu Lys Glu Gly Lys Thr Leu Gly Ala Ile Gln Ile Ile Ile Gly
65 70 75 80

Leu Ala His Ile Gly Leu Gly Ser Ile Met Ala Thr Val Leu Val Gly
85 90 95

Glu Tyr Leu Ser Ile Ser Phe Tyr Gly Gly Phe Pro Phe Trp Gly Gly
100 105 110

Leu Trp Phe Ile Ile Ser Gly Ser Leu Ser Val Ala Ala Glu Asn Gln
115 120 125

Pro Tyr Ser Tyr Cys Leu Leu Ser Gly Ser Leu Gly Leu Asn Ile Val
130 135 140

Ser Ala Ile Cys Ser Ala Val Gly Val Ile Leu Phe Ile Thr Asp Leu
145 150 155 160

Ser Ile Pro His Pro Tyr Ala Tyr Pro Asp Tyr Tyr Pro Tyr Ala Trp
165 170 175

Gly Val Asn Pro Gly Met Ala Ile Ser Gly Val Leu Leu Val Phe Cys
180 185 190

Leu Leu Glu Phe Gly Ile Ala Cys Ala Ser Ser His Phe Gly Cys Gln
195 200 205

Leu Val Cys Cys Gln Ser Ser Asn Val Ser Val Ile Tyr Pro Asn Ile
210 215 220

Tyr Ala Ala Asn Pro Val Ile Thr Pro Glu Pro Val Thr Ser Pro Pro
225 230 235 240

<400> 1679
Leu Val Cys Cys Gln Ser Ser Asn Val Ser Val Ile Tyr Pro Asn Ile

